



STIC Search Report

Biotech-Chem Library

STIC Database Tracking Number: 157955

TO: Jegatheesan Seharaseyon
Art Unit: 1647
Location: REM-4C61/4C70
Serial Number: 09/658677

Thursday, July 07, 2005

From: Beverly Shears
Location: Biotech-Chem Library
REM 1A54
Phone: 571-272-2528
beverly.shears@uspto.gov

Search Notes

Protein Sequence Searches – February 2005

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension **.rup**) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (uniPARC) at:

<http://www.pir.uniprot.org/database/archive.shtml>

If you have any questions regarding this information or your results, please contact any STIC searcher.



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STIC-Biotech/ChemLib

157955

LB

From: Seharaseyon, Jegatheesan
Sent: Thursday, June 30, 2005 9:27 AM
To: STIC-Biotech/ChemLib
Subject: RE:09/658677

CRFE

Hi,

Please search SEQ ID NO: 2, 15 and 18 of 09/658,677
in the interference and commercial databases.

Thanks.
Seyon.

2 - 392 AM
15 - 392 AM
18 - 383 AM

J. Seharaseyon
Art Unit 1647
Remsen 4C61
Mailbox 4C70
Phone: (571)-272-0892
Fax: (571)-273-0892

STAFF USE ONLY

Searcher: _____
Searcher Phone: 2-_____
Date Searcher Picked up: _____
Date Completed: _____
Searcher Prep/Rev. Time: _____
Online Time: _____

Type of Search

NA#: _____ AA#: _____
Interference: _____ SPDI: _____
S/L: _____ Oligomer: _____
Encode/Transl: _____
Structure#: _____ Text: _____
Inventor: _____ Litigation: _____

Vendors and cost where applicable

STN: _____
DIALOG: _____
QUESTEL/ORBIT: _____
LEXIS/NEXIS: _____
SEQUENCE SYSTEM: _____
WWW/Internet: _____
Other(Specify): _____

Date completed: _____

Searcher: Beverly e 2528

Terminal time: _____

Elapsed time: _____

CPU time: _____

Total time: _____

Number of Searches: _____

Number of Databases: _____

Search Site

_____ STIC
_____ CM-1
_____ Pre-S

Type of Search

_____ N.A. Sequence
_____ A.A. Sequence
_____ Structure
_____ Bibliographic

Vendors

_____ IG
_____ STN
_____ Dialog
_____ APS
_____ Geninfo
_____ SDC
_____ DARC/Questel
_____ Other CGN

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: July 1, 2005, 20:52:53 ; Search time 100.435 Seconds
(without alignments)
1509.530 Million cell updates/sec

Title: US-09-658-677-2

Perfect score: 2112

Sequence: 1 MAGIPGLFLFLFLCAVQ.....IKGNVLDREGDVFPLPGSN 392

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 2105692 seqs, 386760381 residues

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : A_Geneseq_16Dec04:*

1: Geneseq1980s:*

2: Geneseq1990s:*

3: Geneseq2000s:*

4: Geneseq2001s:*

5: Geneseq2002s:*

6: Geneseq2003as:*

7: Geneseq2003bs:*

8: Geneseq2004s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	2104	99.6	392	4 AAB48972	Aab48972 Human Zsi
2	2085	98.7	322	4 AAB48973	Aab48973 Human Zsi
3	2044	96.8	383	2 AAY08660	Aay08660 WO9927094
4	2044	96.8	383	2 AAY08657	Aay08657 Human tra
5	2044	96.8	383	2 AAY13390	Aay13390 Amino aci
6	2044	96.8	383	3 AAY88277	Aay88277 Human TAN
7	2044	96.8	383	3 AAY87270	Aay87270 Human sig
8	2044	96.8	383	3 AAY53627	Aay53627 A bone ma
9	2044	96.8	383	3 AAB25618	Aab25618 Protein e
10	2044	96.8	383	3 AAB25592	Aab25592 Protein e
11	2044	96.8	383	3 ADC78573	Adc78573 Human PRO
12	2044	96.8	383	4 AAB80258	Aab80258 Human PRO
13	2044	96.8	383	4 AAB48974	Aab48974 Human Zsi
14	2044	96.8	383	4 AAB29048	Aab29048 Human PRO
15	2044	96.8	383	6 AAB58424	Aab58424 Human PRO
16	2044	96.8	383	6 ABU71636	Abu71636 Human PRO
17	2044	96.8	383	6 ABU87972	Abu87972 Novel hum
18	2044	96.8	383	6 ABU84287	Abu84287 Human sec
19	2044	96.8	383	6 ABR65161	Abr65161 Human sec
20	2044	96.8	383	6 ABR5551	Abr5551 Human sec
21	2044	96.8	383	6 ABU99491	Abu99491 Human sec
22	2044	96.8	383	6 ABU82730	Abu82730 Human PRO
23	2044	96.8	383	6 ABU99851	Abu99851 Novel hum
24	2044	96.8	383	6 ABU71491	Abu71491 Human PRO
25	2044	96.8	383	6 ABR68100	Abr68100 Human sec

26	2044	96.8	383	6 ABU96153	Abu96153 Novel hum
27	2044	96.8	383	6 ABU92584	Abu92584 Human sec
28	2044	96.8	383	6 ABO08661	Abo08661 Human sec
29	2044	96.8	383	6 ABO02713	Abo02713 Human sec
30	2044	96.8	383	6 ABR74867	Abr74867 Human sec
31	2044	96.8	383	6 ABR94629	Abr94629 Human sec
32	2044	96.8	383	6 ABR85602	Abr85602 Human PRO
33	2044	96.8	383	6 ABU98762	Abu98762 Novel hum
34	2044	96.8	383	6 ABU97977	Abu97977 Novel hum
35	2044	96.8	383	6 ABU91683	Abu91683 Novel hum
36	2044	96.8	383	6 ABU71937	Abu71937 Human sec
37	2044	96.8	383	6 ABU89376	Abu89376 Human PRO
38	2044	96.8	383	6 ABU86217	Abu86217 Human sec
39	2044	96.8	383	6 ABU67430	Abu67430 Human sec
40	2044	96.8	383	6 ABU80458	Abu80458 Human PRO
41	2044	96.8	383	6 ABO01820	Abo01820 Novel hum
42	2044	96.8	383	6 ABR99376	Abr99376 Human sec
43	2044	96.8	383	6 ABR98766	Abr98766 Human sec
44	2044	96.8	383	6 ABO16289	Abo16289 Human sec
45	2044	96.8	383	6 ABR92189	Abr92189 Human sec

ALIGNMENTS

RESULT 1
AAB48972
ID AAB48972 standard; protein; 392 AA.
XX
AC AAB48972;
XX
AC
DT 27-MAR-2001 (first entry).
XX
DE Human Zsig13 variant #1, SEQ ID NO:2.
XX
KW Human Zsig13; serine protease; chromosome 11q22.1; elastase homologue;
KW glutamyl endopeptidase homologue; factor X homologue; trypsin homologue;
KW trypsinogen homologue; mast cell protease homologue;
KW collagenase homologue; protein degradation; food processing; brewing;
KW alcohol production; laundry detergent component.
XX
OS Homo sapiens.
XX
PN US6153420-A.
XX
PD 28-NOV-2000.
XX
PF 04-MAY-1998; 98US-00072384.
XX
PR 24-APR-1997; 97US-0044185P.
PR 17-APR-1998; 98US-00062142.
PA (ZYMO) ZYMOGENETICS INC.
XX
PI Sheppard PO;
XX
DR WPI; 2001-060090/07.
DR N-PSDB; AAC91782.
XX
PT New isolated serine protease (designated Zsig13), useful in industrial processes to degrade unwanted proteins or alter the characteristics of protein-containing composition, as well as in industrial applications (e.g. brewing).
XX
PS Claim 1; Col 25-28; 26pp; English.
XX
CC The invention relates to human Zsig13 proteins (AAB48972-B48974), and to DNA encoding them (AAC91782-C91784). The invention also relates to expression vectors and host cells comprising a human Zsig13 DNA, and the recombinant production of a human Zsig13 protein or its precursor. Zsig13 is a serine protease, and has significant homology to Bacillus licheniformis glutamyl endopeptidase, human clotting factor X, human elastase, rat mast cell protease, Streptomyces griseus trypsin, bovine

CC trypsinogen, and Hypoderma lineatum collagenase. The gene encoding human
CC Zsig13 is located on chromosome 11q22.1. Zsig13 is useful in industrial
CC processes to degrade unwanted proteins or alter the characteristics of
CC protein-containing compositions. It may also be used in industrial
CC applications in which proteases are utilised, including food processing,
CC brewing and alcohol production, and as a component of a laundry
CC detergent. The present sequence represents a human Zsig13 variant
XX
SQ Sequence 392 AA;

Query Match 99.6%; Score 2104; DB 4; Length 392;
Best Local Similarity 100.0%; Pred. No. 4.6e-146; Mismatches 0; Indels 0; Gaps 0;
Matches 392; Conservative 0;

Qy 1 MAGIPGLLFLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

Qy 61 VSSSCGPQCHKGTPPTYKEAKQYLSYETLYANGSRTEXQVGIYILSSSGDGAXXRDSGS 120
Db 61 VSSSCGPQCHKGTPPTYKEAKQYLSYETLYANGSRTEXQVGIYILSSSGDGAXXRDSGS 120

Qy 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180

Qy 181 KTYVGTQKLRVGLFKPKFQKGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240
Db 181 KTYVGTQKLRVGLFKPKFQKGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240

Qy 241 IGMVDYALLENLKKPHKPKFMKIGVSPAPKQLPGRIHFGSGYDNDPGLNLYRFCVDKDE 300
Db 241 IGMVDYALLENLKKPHKPKFMKIGVSPAPKQLPGRIHFGSGYDNDPGLNLYRFCVDKDE 300

Qy 301 TYDLLYQCCDAQPGASGYGVYVWMKROOKWERKIIIGFSGHQMVMNGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAQPGASGYGVYVWMKROOKWERKIIIGFSGHQMVMNGSPQDFNVAVR 360

Qy 361 ITPLKYAQICYWIKGNLYDCREGDTVFLPGSN 392
Db 361 ITPLKYAQICYWIKGNLYDCREGDTVFLPGSN 392

RESULT 2
AAB48973
ID AAB48973 standard; protein; 392 AA.
XX
AC AAB48973;
XX
DT 27-MAR-2001 (first entry)
XX
DE Human Zsig13 variant #2, SEQ ID NO:15.
XX
KW Human Zsig13; serine protease; chromosome 11q22.1; elastase homologue;
KW glutamyl endopeptidase homologue; factor X homologue; trypsin homologue;
KW trypsinogen homologue; mast cell protease homologue;
KW collagenase homologue; protein degradation; food processing; brewing;
KW alcohol production; laundry detergent component.
XX
OS Homo sapiens.
XX
XX US6153420-A.
XX
XX 28-NOV-2000.
XX
XX 04-MAY-1998; 98US-00072384.
XX
XX 24-APR-1997; 97US-0044185P.
XX
XX 17-APR-1998; 98US-00062142.
XX
XX (ZYMO) ZYMOGENETICS INC.
XX
XX Sheppard PO;

XX WPI; 2001-060090/07.
DR N-PSDB; AAC91783.
XX
PT New isolated serine protease (designated Zsig13), useful in industrial
PT processes to degrade unwanted proteins or alter the characteristics of
PT protein-containing composition, as well as in industrial applications
PT (e.g. brewing).
XX
PS Claim 1; Col 35-38; 26pp; English.
XX
XX The invention relates to human Zsig13 proteins (AAB48972-B48974), and to
CC DNA encoding them (AAC91782-C91784). The invention also relates to
CC expression vectors and host cells comprising a human Zsig13 DNA, and the
CC recombinant production of a human Zsig13 protein or its precursor. Zsig13
CC is a serine protease, and has significant homology to Bacillus
CC licheniformis glutamyl endopeptidase, human clotting factor X, human
CC elastase, rat mast cell protease, Streptomyces griseus trypsin, bovine
CC trypsinogen, and Hypoderma lineatum collagenase. The gene encoding human
CC Zsig13 is located on chromosome 11q22.1. Zsig13 is useful in industrial
CC processes to degrade unwanted proteins or alter the characteristics of
CC protein-containing compositions. It may also be used in industrial
CC applications in which proteases are utilised, including food processing,
CC brewing and alcohol production, and as a component of a laundry
CC detergent. The present sequence represents a human Zsig13 variant
XX
SQ Sequence 392 AA;

Query Match 98.7%; Score 2085; DB 4; Length 392;
Best Local Similarity 98.2%; Pred. No. 1.1e-144;
Matches 385; Conservative 1; Mismatches 6; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

Qy 61 VSSSCGPQCHKGTPPTYKEAKQYLSYETLYANGSRTEXQVGIYILSSSGDGAXXRDSGS 120
Db 61 VSSSCGPQCHKGTPPTYKEAKQYLSYETLYANGSRTEXQVGIYILSSSGDGAXXRDSGS 120

Qy 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180

Qy 181 KTYVGTQKLRVGLFKPKFQKGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240
Db 181 KTYVGTQKLRVGLFKPKFQKGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240

Qy 241 IGMVDYALLENLKKPHKPKFMKIGVSPAPKQLPGRIHFGSGYDNDPGLNLYRFCVDKDE 300
Db 241 IGMVDYALLENLKKPHKPKFMKIGVSPAPKQLPGRIHFGSGYDNDPGLNLYRFCVDKDE 300

Qy 301 TYDLLYQCCDAQPGASGYGVYVWMKROOKWERKIIIGFSGHQMVMNGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAQPGASGYGVYVWMKROOKWERKIIIGFSGHQMVMNGSPQDFNVAVR 360

Qy 361 ITPLKYAQICYWIKGNLYDCREGDTVFLPGSN 392
Db 361 ITPLKYAQICYWIKGNLYDCREGDTVFLPGSN 392

RESULT 3
AAY08660
ID AAY08660 standard; protein; 383 AA.
XX
AC AAY08660;
XX
DT 09-AUG-1999 (first entry)
XX
XX W09927094 Seq ID 12.
XX
KW Transmembrane domain; human; nutrition; cytokine; cell differentiation;
KW immune stimulation; immune suppression; haematopoiesis; activin;

QY 181 KTVYKGTQKLRVGLKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTTHVTPKGIKGNAND 240
Db 181 KTVYKGTQKLRVGLKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTTHVTPKGIKGNAND 240
QY 241 IGMDYDVALLELKKPKRKFEMKIGVSPAKQLPGGRIFHSGYDNDPFGNLVYRFCDDVKDE 300
Db 241 IGMDYDVALLELKKPKRKFEMKIGVSPAKQLPGGRIFHSGYDNDPFGNLVYRFCDDVKDE 300
QY 301 TYDLLYQCCDAQPGASGYGVYVVMWKRQOQKWERKIIGIFSGHQMVMGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAQPGASGYGVYVVMWKRQOQKWERKIIGIFSGHQMVMGSPQDFNVAVR 360
QY 361 ITPKVAQICWIKGNVLDREG 383
Db 361 ITPKVAQICWIKGNVLDREG 383
RESULT 5
ID AAY13390 standard; protein; 383 AA.
XX AC AAY13390;
XX AC
XX DT 25-JUN-1999 (first entry)
XX XX
XX DE Amino acid sequence of protein PRO307.
XX XX
XX KW Secreted protein; transmembrane protein; human; enterocolitis;
XX KW Zollinger-Ellison syndrome; gastrointestinal ulceration;
XX KW congenital microvillus atrophy; skin disease; cell growth;
XX KW abnormal keratinocyte differentiation; psoriasis; epithelial cancer;
XX KW Parkinson's disease; Alzheimer's disease; ALS; neuropathy; fibromodulin;
XX KW dermal scarring; Usher Syndrome; Atrophia areata; anti-thrombotic;
XX KW wound healing; tissue repair.
XX OS Homo sapiens.
XX XX
XX PN WO9914328-A2..
XX PD
XX PD 25-MAR-1999.
XX PF 16-SEP-1998; 98WO-US019330.
XX PR 17-SEP-1997; 97US-0059113P.
XX PR 17-SEP-1997; 97US-0059115P.
XX PR 17-SEP-1997; 97US-0059117P.
XX PR 17-SEP-1997; 97US-0059119P.
XX PR 17-SEP-1997; 97US-0059121P.
XX PR 17-SEP-1997; 97US-0059122P.
XX PR 17-SEP-1997; 97US-0059184P.
XX PR 18-SEP-1997; 97US-0059263P.
XX PR 18-SEP-1997; 97US-0059266P.
XX PR 18-SEP-1997; 97US-0059266P.
XX PR 17-OCT-1997; 97US-0062125P.
XX PR 17-OCT-1997; 97US-0062285P.
XX PR 17-OCT-1997; 97US-0062287P.
XX PR 21-OCT-1997; 97US-0063486P.
XX PR 24-OCT-1997; 97US-0062814P.
XX PR 24-OCT-1997; 97US-0062816P.
XX PR 24-OCT-1997; 97US-0063045P.
XX PR 24-OCT-1997; 97US-0063120P.
XX PR 24-OCT-1997; 97US-0063121P.
XX PR 24-OCT-1997; 97US-0063127P.
XX PR 24-OCT-1997; 97US-0063128P.
XX PR 27-OCT-1997; 97US-0063327P.
XX PR 27-OCT-1997; 97US-0063329P.
XX PR 28-OCT-1997; 97US-0063341P.
XX PR 28-OCT-1997; 97US-0063342P.
XX PR 28-OCT-1997; 97US-0063542P.
XX PR 28-OCT-1997; 97US-0063549P.
XX PR 28-OCT-1997; 97US-0063550P.
XX PR 28-OCT-1997; 97US-0063564P.
XX PR 28-OCT-1997; 97US-0063435P.
XX PR 29-OCT-1997; 97US-0063704P.

PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
XX (GETH) GENENTECH INC.
XX PA
XX Wood WI, Gurney AL, Goddard A, Pennica D, Chen J, Yuan J;
XX PI
XX XX
XX DR WPI; 1999-229533/19.
XX DR N-PSDB; AAX52261.
XX XX
XX PT New isolated human genes and polypeptides used in, e.g. treatment of
XX PT gastrointestinal ulceration.
XX XX
XX PS Claim 12; Fig 96; 320pp; English.
XX CC
XX CC AAY13344-403 represent secreted and transmembrane human proteins. The
XX CC cDNA sequences are obtained from cDNA libraries, prepared from fetal
XX CC lung, fetal kidney, fetal brain, fetal liver and fetal retina. The
XX CC encoded polypeptides have specific uses based on their homology to known
XX CC polypeptides, e.g. PRO211 and PRO217 can be used for disorders associated
XX CC with the preservation and maintenance of gastrointestinal mucosa and the
XX CC repair of acute and chronic mucosal lesions (e.g. enterocolitis,
XX CC Zollinger-Ellison syndrome, gastrointestinal ulceration and congenital
XX CC microvillus atrophy), skin diseases associated with abnormal keratinocyte
XX CC differentiation (e.g. psoriasis, epithelial cancers such as lung squamous
XX CC cell carcinoma of the vulva and gliomas), potent effects on cell growth
XX CC and development, diseases related to growth or survival of nerve cells
XX CC including Parkinson's disease, Alzheimer's disease, ALS, neuropathies or
XX CC cancer. PRO265 can be used as for fibromodulin, e.g. for reducing dermal
XX CC scarring. PRO264 can be used as a target for anti-tumor drugs. PRO533 may
XX CC be used in the treatment of Usher Syndrome or Atrophia areata; PRO269 can
XX CC be used as an anti-thrombotic agent; PRO287 polypeptides and portions may
XX CC have therapeutic applications in wound healing and tissue repair; PRO317
XX CC can be used for treating problems of the kidney, uterus, endometrium,
XX CC blood vessels, or related tissue, e.g. in the heart of genital tract
XX XX
XX SQ Sequence 383 AA;
Query Match 96.8%; Score 2044; DB 2; Length 383;
Best Local Similarity 98.4%; Pred. No. 1.1e-141;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;
QY 1 MAGIFGLLFLFFLLCAVQGVSPYSAPKWPAYRLPVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIFGLLFLFFLLCAVQGVSPYSAPKWPAYRLPVLPQSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPGCHGKTPLPTTYKEAKQYLSYETLYANGSRTXQVIGIYILSSGDCAXXRDSSG 120
Db 61 VSSSCGPGCHGKTPLPTTYKEAKQYLSYETLYANGSRTXQVIGIYILSSGDCAXXRDSSG 120
QY 121 SGKSRKRFQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
Db 121 SGKSRKRFQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
QY 181 KTVYKGTQKLRVGLKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTTHVTPKGIKGNAND 240

Db 181 KTVVGTQKLRVGLFKPKFGKGRGANDSTSAPEQMKFQWIRKTHVPKGIKGNAND 240
 Qy 241 IGMDDYALLELKKPKHKKFMKIGVSPAKQLPGRIHFGSYNDNRPNLVYRFDVKDE 300
 Db 241 IGMDDYALLELKKPKHKKFMKIGVSPAKQLPGRIHFGSYNDNRPNLVYRFDVKDE 300
 Qy 301 TYDLLVQCCDAQPGASGYGVYVVMWKRQKQKWKRIIGIFSGHQWVDMNGSPQDFNVAVR 360
 Db 301 TYDLLVQCCDAQPGASGYGVYVVMWKRQKQKWKRIIGIFSGHQWVDMNGSPQDFNVAVR 360
 Qy 361 ITPLKYAQCICWIKGNLYDCREG 383
 Db 361 ITPLKYAQCICWIKGNLYDCREG 383

RESULT 6

AAV88277
 ID AAY88277 standard; protein; 383 AA.

AC AAY88277;

XX 16-OCT-2000 (first entry)

DT Human TANGO 186 protein.

DE TANGO 180; TANGO 181; TANGO 182; TANGO 183; TANGO 184; TANGO 185;

XX TANGO 186; TANGO 188; TANGO 189; TANGO 215; TANGO 187; human; murine;

XX secreted protein; transmembrane protein; gene therapy; vaccine;

XX diagnosis; treatment; detection.

XX Homo sapiens.

XX WO200018904-A2.

XX 06-APR-2000.

XX 30-SEP-1999; 99WO-US022817.

XX 30-SEP-1998; 98US-00164220.

PR 02-OCT-1998; 98US-00164169.

XX (MILL-) MILLENNIUM BIOTHERAPEUTICS INC.

XX Barnes TM;

XX WPI; 2000-293144/25.

DR N-PSDB; AAA39945, AAA39946.

XX Isolated nucleic acids encoding TANGO polypeptides useful for preventing,

PT diagnosing and treating diseases associated with inappropriate protein

PT expression.

XX Claim 9; Fig 13; 249pp; English.

PS This invention describes novel human and murine nucleic acids encoding

CC TANGO polypeptides (which are either wholly secreted or transmembrane

CC proteins) which can be used for gene therapy and/or vaccination. The

CC peptides are designated TANGO 180 to TANGO 189 and TANGO 215. The nucleic

CC acids may be used to produce TANGO 180 to TANGO 189 and TANGO 215

CC polypeptides according to standard recombinant DNA methodologies. They

CC may also be used to detect and identify the presence of TANGO nucleic

CC acids in a sample and therefore identify or diagnose diseases associated

CC with inappropriate TANGO expression (e.g. diseases related to over or

CC under expression of the polypeptides or the expression of inactive

CC polypeptides). The nucleic acids and the polypeptides they encode may be

CC used according to standard gene therapy protocols, to treat diseases

CC associated with inappropriate TANGO expression by supplementing a

CC patients own production of the polypeptide of to rectify mutations that

CC may result in expression of an abnormally active polypeptide. The

CC polypeptides may also be used to identify and produce agonists and

CC antagonists of TANGO expression and activity which may be used to

CC modulate TANGO related processes and diseases. The polypeptides are

CC particularly useful for use as antigens for producing antibodies to TANGO

CC proteins which may be used for inhibiting the activity of TANGO proteins.

CC They may also be used to detect and quantify the presence of TANGO

CC proteins in samples and therefore identify patients in whom the protein

CC is over- or under-expressed. This sequence represents the human TANGO 186

CC protein described in the method of the invention

XX SQ Sequence 383 AA;

Query Match 96.8%; Score 2044; DB 3; Length 383;

Best Local Similarity 98.4%; Pred. No. 1.1e-141;

Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFLCAVGVSPYSAPWKPPTWYRLPVVLPSTLNLAKEPFGASAKLE 60

Db 1 MAGIPGLLFLFLCAVGVSPYSAPWKPPTWYRLPVVLPSTLNLAKEPFGASAKLE 60

Qy 61 VSSSCGPGCHKGTPLPTTKEAKQYLSYETLYANGSRTEXQVGIYIILSSGSGAXRDSGS 120

Db 61 VSSSCGPGCHKGTPLPTTKEAKQYLSYETLYANGSRTEXQVGIYIILSSGSGAXRDSGS 120

Qy 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLTCTGTTLVAEXHVLTAHCIDHG 180

Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLTCTGTTLVAEXHVLTAHCIDHG 180

Qy 181 KTVVGTQKLRVGLFKPKFGKGRGANDSTSAPEQMKFQWIRKTHVPKGIKGNAND 240

Db 181 KTVVGTQKLRVGLFKPKFGKGRGANDSTSAPEQMKFQWIRKTHVPKGIKGNAND 240

Qy 241 IGMDDYALLELKKPKHKKFMKIGVSPAKQLPGRIHFGSYNDNRPNLVYRFDVKDE 300

Db 241 IGMDDYALLELKKPKHKKFMKIGVSPAKQLPGRIHFGSYNDNRPNLVYRFDVKDE 300

Qy 301 TYDLLVQCCDAQPGASGYGVYVVMWKRQKQKWKRIIGIFSGHQWVDMNGSPQDFNVAVR 360

Db 301 TYDLLVQCCDAQPGASGYGVYVVMWKRQKQKWKRIIGIFSGHQWVDMNGSPQDFNVAVR 360

Qy 361 ITPLKYAQCICWIKGNLYDCREG 383

Db 361 ITPLKYAQCICWIKGNLYDCREG 383

RESULT 7

AAV87270

ID AAY87270 standard; protein; 383 AA.

XX AAY87270;

XX 11-MAY-2000 (first entry)

DT Human signal peptide containing protein HSP-47 SEQ ID NO:47.

DE Human; signal peptide-containing protein; HSP-47; diagnosis; cancer;

XX inflammation; cardiovascular disease; anticancer; anti-inflammatory;

XX antimicrobial; neurotropic; neuroprotective; cardiovascular; hepatotropic;

XX antitastmatic; gene therapy; cell proliferation; neurological disorder;

XX reproductive disorder; developmental disorder; arteriosclerosis;

XX cirrhosis; psoriasis; acquired immune deficiency syndrome; anaemia;

XX asthma; Crohn's disease; infection; Alzheimer's disease; schizophrenia;

XX Parkinson's disease; Huntington's disease; ovulatory defect;

XX muscular dystrophy.

XX Homo sapiens.

OS WO200000610-A2.

PN 06-JAN-2000.

PD 25-JUN-1999; 99WO-US014484.

XX 26-JUN-1998; 98US-0090762P.

PR 31-JUL-1998; 98US-0094983P.

PR 01-OCT-1998; 98US-0102686P.

CC and ulcers, to induce cartilage and/or bone growth in circumstances where
 CC bone is not normally formed and thus have an application in healing bone
 CC fractures and cartilage damage or defects, prophylactic use in fracture
 CC reduction and also in the improved fixation of artificial joints
 XX SQ Sequence 383 AA;

Query Match 96.8%; Score 2044; DB 3; Length 383;
 Best Local Similarity 98.4%; Pred. No. 1.1e-141;
 Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAGVQSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
 DB 1 MAGIPGLLFLFLCAGVQSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPGQCHKTPLPTYPEAKQVLSYETLYANGSRTEKQVGIYILSSSGDGAAXRDSGS 120
 DB 61 VSSSCGPGQCHKTPLPTYPEAKQVLSYETLYANGSRTEKQVGIYILSSSGDGAAXRDSGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
 DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180

QY 181 KTVVGTQKLRVGLFKPKFDCGGRANDSTSAMPEQMKFQWIRKTRTHVPKGIKGNAND 240
 DB 181 KTVVGTQKLRVGLFKPKFDCGGRANDSTSAMPEQMKFQWIRKTRTHVPKGIKGNAND 240

QY 241 IGMVDYDYLLELKKPKHKKFPMKIGVSPAPKQLPGRHFGSYNDPRLNLYRFDCKDE 300
 DB 241 IGMVDYDYLLELKKPKHKKFPMKIGVSPAPKQLPGRHFGSYNDPRLNLYRFDCKDE 300

QY 301 TYDLLYQCCDAQPGASGYGVYVMWKRQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
 DB 301 TYDLLYQCCDAQPGASGYGVYVMWKRQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360

QY 361 ITPLKYAQICYWKGNLYDCREG 383
 DB 361 ITPLKYAQICYWKGNLYDCREG 383

RESULT 9
 AAB25618
 ID AAB25618 standard; protein; 383 AA.
 XX AAB25618;
 AC AAB25618;
 XX 21-NOV-2000 (first entry)
 XX Protein encoded by human secreted protein gene #10.
 DE Secreted protein; immunosuppressant; anti-inflammatory; antiarthritic;
 XX antirheumatic, dermatological; antiproliferative; antiarteriosclerotic;
 KW anticancer; vulnary; antiviral; antibacterial; antifungal;
 KW immune disorder; Addison's disease; rheumatoid arthritis; dermatitis;
 KW multiple sclerosis; inflammatory disorder; inflammatory bowel disease;
 KW Crohn's disease; nephritis; hyperproliferative disorder;
 KW cardiovascular disorder; coronary arteriosclerosis; myocarditis; cancer;
 KW melanoma; lymphoma; wound healing; human; chromosome 12.
 XX Homo sapiens.
 OS
 XX WO200029435-A1.
 PN
 XX 25-MAY-2000.
 PD
 XX 27-OCT-1999; 99WO-US025031.
 PF
 XX 28-OCT-1998; 98US-0105971P.
 PR
 XX (HUMA-) HUMAN GENOME SCI INC.
 PA
 XX Ni J, Ruben SM, Olsen HS, Young PE, Kenny JJ, Moore PA, Wei Y;
 PI Greene JW;

XX WPI; 2000-387742/33.
 XX Isolated nucleic acid molecules encoding human secreted proteins are used
 PT for the prevention, amelioration and treatment of autoimmune,
 PT inflammatory, hyperproliferative and cardiovascular disorders, cancer,
 PT wounds, and infectious diseases.

PS Disclosure; Page 169; 803pp; English.
 XX The present invention relates to 12 secreted human proteins and the
 CC nucleotide sequences encoding them. The polynucleotide sequences given in
 CC AAB0606-A80623 encode the 12 secreted protein sequences given in
 CC AAB25576-B25593. The human secreted proteins have various activities
 CC dependent on the tissues in which they are expressed. Examples of the
 CC activities of the proteins include: immunosuppressant; anti-inflammatory;
 CC antitumor; antirheumatic, dermatological; antiproliferative;
 CC antiarteriosclerotic; anticancer; vulnary; antiviral; antibacterial;
 CC and antifungal activity. The proteins, polypeptides, agonists and
 CC antagonists may be used to treat prevent and/or diagnose various disease,
 CC disorders and conditions examples of which include: immune disorders e.g.
 CC Addison's disease, rheumatoid arthritis, dermatitis, and multiple
 CC sclerosis; inflammatory disorders e.g. inflammatory bowel disease,
 CC Crohn's disease and nephritis; hyperproliferative disorders such as
 CC paraproteinemia and purpura; cardiovascular disorders e.g. coronary
 CC arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The
 CC proteins and polynucleotide sequences may also be used in wound healing
 CC and the treatment of infectious diseases. The human secreted protein gene
 CC #10 and protein sequences are represented in sequences AAB0615 and
 CC AAB25585. Secreted protein gene #10 is located on chromosome 12.
 CC Sequences AAB25616-B25618 represent alternative secreted protein #10
 CC sequences and AAB0669-A80676 represent genes which are related to the
 CC secreted protein gene#10
 XX Sequence 383 AA;

Query Match 96.8%; Score 2044; DB 3; Length 383;
 Best Local Similarity 98.4%; Pred. No. 1.1e-141;
 Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAGVQSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
 DB 1 MAGIPGLLFLFLCAGVQSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPGQCHKTPLPTYPEAKQVLSYETLYANGSRTEKQVGIYILSSSGDGAAXRDSGS 120
 DB 61 VSSSCGPGQCHKTPLPTYPEAKQVLSYETLYANGSRTEKQVGIYILSSSGDGAAXRDSGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
 DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180

QY 181 KTVVGTQKLRVGLFKPKFDCGGRANDSTSAMPEQMKFQWIRKTRTHVPKGIKGNAND 240
 DB 181 KTVVGTQKLRVGLFKPKFDCGGRANDSTSAMPEQMKFQWIRKTRTHVPKGIKGNAND 240

QY 241 IGMVDYDYLLELKKPKHKKFPMKIGVSPAPKQLPGRHFGSYNDPRLNLYRFDCKDE 300
 DB 241 IGMVDYDYLLELKKPKHKKFPMKIGVSPAPKQLPGRHFGSYNDPRLNLYRFDCKDE 300

QY 301 TYDLLYQCCDAQPGASGYGVYVMWKRQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
 DB 301 TYDLLYQCCDAQPGASGYGVYVMWKRQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360

QY 361 ITPLKYAQICYWKGNLYDCREG 383
 DB 361 ITPLKYAQICYWKGNLYDCREG 383

RESULT 10
 AAB25592
 ID AAB25592 standard; protein; 383 AA.
 XX

AC AAB25592;
 XX 21-NOV-2000 (first entry)
 XX Protein encoded by human secreted protein gene #10 clone HUSQ05.
 XX
 XX Secreted protein; immunosuppressant; anti-inflammatory; antiarthritic;
 KW antirheumatic; dermatological; antiproliferative; antiarteriosclerotic;
 KW anticancer; vulnary; antiviral; antibacterial; antifungal;
 KW immune disorder; Addison's disease; rheumatoid arthritis; dermatitis;
 KW multiple sclerosis; inflammatory disorder; inflammatory bowel disease;
 KW Crohn's disease; nephritis; hyperproliferative disorder;
 KW cardiovascular disorder; coronary arteriosclerosis; myocarditis; cancer;
 KW melanoma; lymphoma; wound healing; human; chromosome 12.
 XX
 XX Homo sapiens.
 XX
 XX W0200029435-A1.
 XX
 XX 25-MAY-2000.
 XX
 XX 27-OCT-1999; 99WO-US025031.
 XX
 XX 28-OCT-1998; 98US-0105971P.
 XX
 XX (HUMA-) HUMAN GENOME SCI INC.
 XX
 XX Ni J, Ruben SM, Olsen HS, Young PE, Kenny JJ, Moore PA, Wei Y;
 PI Greene JM;
 XX
 XX WPI; 2000-387742/33.
 XX
 XX Isolated nucleic acid molecules encoding human secreted proteins are used
 PT for the prevention, amelioration and treatment of autoimmune,
 PT inflammatory, hyperproliferative and cardiovascular disorders, cancer,
 PT wounds, and infectious diseases.
 XX
 XX Claim 1; Page 684-685; 803pp; English.
 XX
 XX The present invention relates to 12 secreted human proteins and the
 CC nucleotide sequences encoding them. The polynucleotide sequences given in
 CC AAB0606-A80623 encode the 12 secreted protein sequences given in
 CC AAB25576-B25593. The human secreted proteins have various activities
 CC dependent on the tissues in which they are expressed. Examples of the
 CC activities of the proteins include: immunosuppressant; anti-inflammatory;
 CC antiarthritic; antirheumatic; dermatological; antiproliferative;
 CC antiarteriosclerotic; anticancer; vulnary; antiviral; antibacterial;
 CC and antifungal activity. The proteins, polypeptides, agonists and
 CC antagonists may be used to treat prevent and/or diagnose various disease,
 CC disorders and conditions examples of which include: immune disorders e.g.
 CC Addison's disease, rheumatoid arthritis, dermatitis, and multiple
 CC sclerosis; inflammatory disorders e.g. inflammatory bowel disease,
 CC Crohn's disease and nephritis; hyperproliferative disorders such as
 CC paraproteinemias and purpura; cardiovascular disorders e.g. coronary
 CC arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The
 CC proteins and polynucleotide sequences may also be used in wound healing
 CC and the treatment of infectious diseases. The human secreted protein gene
 CC #10 and protein sequences are represented in sequences AAB0615 and
 CC AAB25585. Secreted protein gene #10 is located on chromosome 12.
 CC Sequences AAB25616-B25618 represent alternative secreted protein #10
 CC sequences and AAB0669-A80676 represent genes which are related to the
 CC secreted protein gene#10
 XX
 XX Sequence 383 AA;
 Query Match 96.8%; Score 2044; DB 3; Length 383;
 Best Local Similarity 98.4%; Pred. No. 1.1e-141;
 Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;
 QY 1 MAGIPGLLFLFLLCAGVGQSPYSAPWKPWPAYRLPVVLPOSTNLAKPFGAIAKLE 60
 DB 1 MAGIPGLLFLFLLCAGVGQSPYSAPWKPWPAYRLPVVLPOSTNLAKPFGAIAKLE 60

QY 61 VSSCGPQCHKGTPLPTYPEAKQYLSYETLYANGSRTEKQVGIYILSSSGDGAXXRDSGS 120
 DB 61 VSSCGPQCHKGTPLPTYPEAKQYLSYETLYANGSRTEKQVGIYILSSSGDGAXXRDSGS 120
 QY 121 SKSRRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
 DB 121 SKSRRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
 QY 181 KTYVRGTQKLRVGFLLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTVPGWKIGNAND 240
 DB 181 KTYVRGTQKLRVGFLLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTVPGWKIGNAND 240
 QY 241 IGMVDYDYLLELKKPKFKFKMIGVSPAPKQIPLGRIHFGSYDNDPGLNVRFCVDKDE 300
 DB 241 IGMVDYDYLLELKKPKFKFKMIGVSPAPKQIPLGRIHFGSYDNDPGLNVRFCVDKDE 300
 QY 301 TYDLLYQQCDAOPGASGVGYVYRMKROQKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
 DB 301 TYDLLYQQCDAOPGASGVGYVYRMKROQKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
 QY 361 ITPLKYAQICYWIKGNLYLDCREG 383
 DB 361 ITPLKYAQICYWIKGNLYLDCREG 383
 RESULT 11
 ADC78573
 ID ADC78573 standard; protein; 383 AA.
 XX
 AC ADC78573;
 XX
 DT 01-JAN-2004 (first entry)
 DE Human PRO307 protein.
 KW antinflammatory; antiulcer; cytostatic; antipsoriatic; antiparkinsonian;
 KW neurotrophic; neuroprotective; vasotropic; chemotactic; angiogenic;
 KW antiarteriosclerotic; cardiasthmatic; antiarthritic; antirheumatic;
 KW thrombolytic; immunomodulator; enterocolitis; Zollinger-Ellison syndrome;
 KW gastrointestinal ulceration; psoriasis; cancer; Parkinson's disease;
 KW Alzheimer's; ALS; neuropathy; dermal scarring; wound healing;
 KW nerve repair; thrombosis; bone; cartilage formation; angiogenesis;
 KW asthma; rheumatoid arthritis; multiple sclerosis; inflammatory disorder;
 KW atherosclerosis; cardiac injury; infertility; premature aging; AIDS;
 KW diabetes; stroke; gene therapy; transgenic; PRO; human.
 XX
 OS Homo sapiens.
 XX
 XX WO200015796-A2.
 XX
 XX 23-MAR-2000.
 XX
 XX 15-SEP-1999; 99WO-US021090.
 XX
 XX 16-SEP-1998; 98WO-US019330.
 XX
 XX (GETH) GENENTECH INC.
 XX
 XX Chen J, Goddard A, Gurney AL, Hillan K, Pennica D, Wood WI;
 PI Yuan J;
 XX
 XX WPI; 2000-271434/23.
 XX
 XX N-PSDB; ADC78572.
 XX
 XX Novel nucleic acids encoding secreted and transmembrane polypeptides with
 PT homology, e.g. to growth and cancer-associated antigens.
 XX
 XX Claim 12; SEQ ID NO 261; 355pp; English.
 XX
 XX The invention relates to a novel nucleic acid encoding a PRO polypeptide.
 CC The polypeptides and polynucleotides of the invention may be useful as
 CC research tools and as therapeutics for treating enterocolitis, Zollinger-

CC Ellison syndrome, gastrointestinal ulceration, psoriasis, cancer,
 CC Parkinson's disease, Alzheimer's disease, ALS, neuropathies, dermal
 CC scarring and wound healing, nerve repair, thrombosis, bone and/or
 CC cartilage formation, angiogenesis, asthma, rheumatoid arthritis, multiple
 CC sclerosis, inflammatory disorders, atherosclerosis, cardiac injury,
 CC infertility, premature aging, AIDS, diabetes complications and stroke.
 CC The molecules may also be utilised during gene therapy procedures and
 CC transgenic animal production. The current sequence is that of the human
 CC PRO protein of the invention.
 XX
 SQ Sequence 383 AA;

Query Match 96.8%; Score 2044; DB 3; Length 383;
 Best Local Similarity 98.4%; Pred. No. 1.1e-141;
 Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYAPWKPTWPAVRLPVVLPQSTLNLAKEPFGAEAKLE 60
 DB 1 MAGIPGLLFLFLCAVGVSPYAPWKPTWPAVRLPVVLPQSTLNLAKEPFGAEAKLE 60
 QY 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXRDSGS 120
 DB 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXRDSGS 120
 QY 121 SGSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGCTGTLVAEKHVLTAAHCHIDG 180
 DB 121 SGSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGCTGTLVAEKHVLTAAHCHIDG 180
 QY 181 KTYVKGQKLRVGLFKPKFQDGGGRANDSTSAPEQMKFQWIRKTRHVPKGIKGNAND 240
 DB 181 KTYVKGQKLRVGLFKPKFQDGGGRANDSTSAPEQMKFQWIRKTRHVPKGIKGNAND 240
 QY 241 IGMDYDVALLELKKPHKPKMKIGVSPAPKQIPGRIHFGSYNDPRGNLVYRFDVKDE 300
 DB 241 IGMDYDVALLELKKPHKPKMKIGVSPAPKQIPGRIHFGSYNDPRGNLVYRFDVKDE 300
 QY 301 TYDLLYQCCDAQPGASGVYVVRMKRQKQKWKRIIGIFSGHGWDMNGSPQDFNVAVR 360
 DB 301 TYDLLYQCCDAQPGASGVYVVRMKRQKQKWKRIIGIFSGHGWDMNGSPQDFNVAVR 360
 QY 361 ITELKVAQICWIKGNVLDREG 383
 DB 361 ITELKVAQICWIKGNVLDREG 383

RESULT 12

AAB80258

ID AAB80258 standard; protein; 383 AA.

AC AAB80258;

XX

DT 24-APR-2001 (first entry)

DE Human PRO307 protein.

XX

XX Human; PRO; dermatological; antipsoriatic; cytostatic; antiinflammatory;

XX antiParkinsonian neurotropic; neuroprotective; vulnerary; cardiant;

XX antiangiogenic; vasotropic; antiasthmatic; antirheumatic; cancer;

XX antiarthritic; antifertility; antidiabetic; antiviral; diabetes;

XX opthalmological; gene therapy; skin disease; gastrointestinal disorder;

XX ischaemia; inflammation.

OS Homo sapiens.

XX WO200104311-A1.

PN 18-JAN-2001.

PD

XX 22-FEB-2000; 2000WO-US0004414.

XX

XX 07-JUL-1999; 99US-0143048P.

XX 26-JUL-1999; 99US-0145698P.

XX 28-JUL-1999; 99US-0146222P.

XX

XX

XX

XX

XX

XX

XX

XX

PR 08-SEP-1999; 99WO-US020594.
 PR 13-SEP-1999; 99WO-US020944.
 PR 15-SEP-1999; 99WO-US021090.
 PR 15-SEP-1999; 99WO-US021547.
 PR 05-OCT-1999; 99WO-US023089.
 PR 29-NOV-1999; 99WO-US028214.
 PR 30-NOV-1999; 99WO-US028313.
 PR 02-DEC-1999; 99WO-US028564.
 PR 02-DEC-1999; 99WO-US028565.
 PR 16-DEC-1999; 99WO-US030095.
 PR 20-DEC-1999; 99WO-US030911.
 PR 20-DEC-1999; 99WO-US030999.
 PR 05-JAN-2000; 2000WO-US000219.
 XX (GETH) GENENTECH INC.
 XX

PI Ashkenazi AJ, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
 PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
 PI Godowski PJ, Grimaldi CJ, Gurney AL, Hillan KJ, Kljavin IJ;
 PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
 PI Williams PM, Wood WI;
 XX WPI; 2001-081051/09.
 DR N-PSDB; AAF72419.

XX Sixty one nucleic acids encoding PRO polypeptides which are useful in the
 PT treatment of skin diseases (e.g. psoriasis), cancers (e.g. lung squamous
 PT cell carcinoma) and neurodegenerative diseases (e.g. Alzheimer's
 PT disease).
 XX
 PS Claim 1; Fig 96; 393pp; English.

XX The present sequence is one of sixty one novel secreted and transmembrane
 CC PRO polypeptides. The PRO polypeptides are useful for treating skin
 CC diseases (e.g. psoriasis), cancers (e.g. lung squamous cell carcinoma),
 CC gastrointestinal disorders (e.g. enterocolitis), neurodegenerative
 CC diseases (e.g. Alzheimer's disease, Parkinson's disease), wound repair,
 CC cardiovascular disorders (e.g. endometrial bleeding angiogenesis,
 CC ischaemias such as coronary ischaemia, atherosclerosis), inflammatory
 CC disorders (e.g. asthma, rheumatoid arthritis, multiple sclerosis),
 CC infertility, AIDS and diabetes and retinal disorders such as retinitis
 CC pigmentosum. The PRO nucleic acids have applications in molecular
 CC biology, including use as hybridization probes, and in chromosome and
 CC gene mapping
 XX
 SQ Sequence 383 AA;

Query Match 96.8%; Score 2044; DB 4; Length 383;

Best Local Similarity 98.4%; Pred. No. 1.1e-141;

Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYAPWKPTWPAVRLPVVLPQSTLNLAKEPFGAEAKLE 60
 DB 1 MAGIPGLLFLFLCAVGVSPYAPWKPTWPAVRLPVVLPQSTLNLAKEPFGAEAKLE 60
 QY 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXRDSGS 120
 DB 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXRDSGS 120
 QY 121 SGSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGCTGTLVAEKHVLTAAHCHIDG 180
 DB 121 SGSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGCTGTLVAEKHVLTAAHCHIDG 180
 QY 181 KTYVKGQKLRVGLFKPKFQDGGGRANDSTSAPEQMKFQWIRKTRHVPKGIKGNAND 240
 DB 181 KTYVKGQKLRVGLFKPKFQDGGGRANDSTSAPEQMKFQWIRKTRHVPKGIKGNAND 240
 QY 241 IGMDYDVALLELKKPHKPKMKIGVSPAPKQIPGRIHFGSYNDPRGNLVYRFDVKDE 300
 DB 241 IGMDYDVALLELKKPHKPKMKIGVSPAPKQIPGRIHFGSYNDPRGNLVYRFDVKDE 300
 QY 301 TYDLLYQCCDAQPGASGVYVVRMKRQKQKWKRIIGIFSGHGWDMNGSPQDFNVAVR 360

Db	301	TYDLLYQQCDAQPGASGSGVYVVMKRWKQQQKWERKIIGIFSGHQWVDMNGSPQDFNVAVR	360
Qy	361	ITPLKYAICYWIKGNLYDCREG 383	
Db	361	ITPLKYAICYWIKGNLYDCREG 383	
RESULT 13			
AA48974			
ID	AA48974	standard; protein; 383 AA.	
AC	AA48974;		
XX			
DT	27-MAR-2001	(first entry)	
XX			
DE	Human Zsig13 variant #3, SEQ ID NO:18.		
XX			
KW	Human Zsig13; serine protease; chromosome 11q22.1; elastase homologue;		
KW	glutamyl endopeptidase homologue; factor X homologue; trypsin homologue;		
KW	trypsinogen homologue; mast cell protease homologue;		
KW	collagenase homologue; protein degradation; food processing; brewing;		
KW	alcohol production; laundry detergent component.		
XX			
OS	Homo sapiens.		
XX			
FN	US6153420-A.		
XX			
PD	28-NOV-2000.		
XX			
PF	04-MAY-1998;	98US-00072384.	
XX			
PR	24-APR-1997;	97US-0044185P.	
PR	17-APR-1998;	98US-00062142.	
XX			
PA	(ZYMO) ZYMOGENETICS INC.		
XX			
PI	Sheppard PO;		
XX			
DR	WPI; 2001-060090/07.		
DR	N-PSDB; AAC91784.		
XX			
PT	New isolated serine protease (designated Zsig13), useful in industrial		
PT	processes to degrade unwanted proteins or alter the characteristics of		
PT	protein-containing composition, as well as in industrial applications		
PT	(e.g. brewing).		
XX			
PS	Claim 1; Col 41-44; 26pp; English.		
XX			
CC	The invention relates to human Zsig13 proteins (AAB48972-548974), and to		
CC	DNA encoding them (AAC91782-C91784). The invention also relates to		
CC	expression vectors and host cells comprising a human Zsig13 DNA, and the		
CC	recombinant production of a human Zsig13 protein or its precursor. Zsig13		
CC	is a serine protease, and has significant homology to Bacillus		
CC	licheniformis glutamyl endopeptidase, human clotting factor X, human		
CC	elastase, rat mast cell protease, Streptomyces griseus trypsin, bovine		
CC	trypsinogen, and Hypoderma lineatum collagenase. The gene encoding human		
CC	Zsig13 is located on chromosome 11q22.1. Zsig13 is useful in industrial		
CC	processes to degrade unwanted proteins or alter the characteristics of		
CC	protein-containing compositions. It may also be used in industrial		
CC	applications in which proteases are utilised, including food processing,		
CC	brewing and alcohol production, and as a component of a laundry		
CC	detergent. The present sequence represents a human Zsig13 variant		
XX			
SQ	Sequence 383 AA;		
Query Match			
Best Local Similarity		96.8%;	Score 2044; DB 4; Length 383;
Matches 377; Conservative		1;	Mismatches 5; Indels 0; Gaps 0;
Qy	1	MAGIPGLLFLFLLCAGVGQVSPYAPWKTWPAYRLPVVLPQSTLNLAKPFGAEAKLE	60
Db	1	MAGIPGLLFLFLLCAGVGQVSPYAPWKTWPAYRLPVVLPQSTLNLAKPFGAEAKLE	60

Qy	61	VSSCGPQCHKTPTPTYPEAKQYLSYETLYANGSRTXQVCIYILSSSGDGAXXRDSGS	120
Db	61	VSSCGPQCHKTPTPTYPEAKQYLSYETLYANGSRTTETQVGIYILSSSGDGQAHRDSGS	120
Qy	121	SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEXHVLTAACHIDG	180
Db	121	SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG	180
Qy	181	KTYVKGTKLVRGFLKPKFKDGGRGANDSTAMPQMKFQWIRVKRTHVPKGIKGNAND	240
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KW	dog; cat; pig; goat; rabbit; tumour necrosis factor alpha; TNF-alpha;		
KW	blood; chondrocyte cell; cell proliferation; cell differentiation; colon;		
KW	adrenal; lung; breast; prostate; rectum; cervix; liver; genetic disorder.		
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OS	Homo sapiens.		
XX			
PN	WO200168848-A2.		
XX			
PD	20-SEP-2001.		
XX			
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PR 20-DEC-2000; 2000WO-US034956.
XX
XX (GETH ) GENENTECH INC.
XX
XX Baker KP, Chen J, Deanoyers L, Goddard A, Godowski PJ, Gurney AL;
XX Pan J, Smith V, Watanabe CK, Wood WI, Zhang Z;
XX
XX WPI; 2001-602746/68.
XX N-PSDB; AAS45949.
XX
XX Novel nucleic acids encoding PRO polypeptides, used to diagnose the
XX presence of tumors, such as prostate and breast tumors, in mammals and to
XX screen for modulators of the compounds.
XX
XX Claim 11; Fig 50; 774pp; English.
XX
XX Sequences AAU29024-AAU29328 represent PRO polypeptides of the invention.
XX The PRO polypeptides and their associated nucleic acids can be used to
XX detect the presence of a tumour in a mammal by comparing the level of
XX expression of a PRO polypeptide in a test sample of cells from the animal
XX and a control sample of normal cells, whereby a higher level of
XX expression in the test sample indicates the presence of a tumour in the
XX mammal. Mammals include dogs, cats, cattle, horses, sheep, pigs, goats
XX and rabbits but are preferably human. The polypeptides can be used to
XX stimulate tumour necrosis factor (TNF) alpha release from human blood,
XX when contacted with it. A specific polypeptide can be used to stimulate
XX the proliferation or differentiation of chondrocyte cells. The PRO
XX proteins can be used to determine the presence of tumours and also
XX susceptibility to tumour development, particularly adrenal, lung, colon,
XX breast, prostate, rectal, cervical, or liver tumours, in mammalian
XX subjects. The oligonucleotide probes specific for the PRO nucleic acids
XX can be used for genetic analysis of individuals with genetic disorders
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XX antibody-dependent enzyme mediated prodrug therapy.
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OM protein - protein search, using sw model

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Post-processing: Minimum Match 0%

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43 114 5.4 254 1 US-08-330-978-3 Sequence 3, Appl
44 114 5.4 254 1 US-08-474-042-3 Sequence 3, Appl
45 114 5.4 254 1 US-08-484-558-3 Sequence 3, Appl

ALIGNMENTS

RESULT 1

US-09-072-384-2
; Sequence 2, Application US/09072384
; Patent No. 6153420
; GENERAL INFORMATION:
; APPLICANT: Sheppard, Paul O.
; TITLE OF INVENTION: SERINE PROTEASE POLYPEPTIDES
; TITLE OF INVENTION: AND MATERIALS AND METHODS FOR MAKING THEM
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: ZymoGenetics, Inc.
; STREET: 1201 Eastlake Avenue East
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98102

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/072,384
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Parker, Gary E
REGISTRATION NUMBER: 31,648
REFERENCE/DOCKET NUMBER: 97-16C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-442-6673
TELEFAX: 206-442-6678
TELEX:
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 392 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
FRAGMENT TYPE: internal
FEATURE:
NAME/KEY: Signal Sequence
LOCATION: 1...19
OTHER INFORMATION:
US-09-072-384-2

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Query Match          99.6%; Score 2104; DB 3; Length 392;
Best Local Similarity 100.0%; Pred. No. 5.2e-218;
Matches 392; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
    |||||
Db 1 MAGIPGLLFLFLCAVGVSPYSAPWKPPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
    |||||

QY 61 VSSSCGPOCHKGTPPTYKEAKQYLSYETLYANGSRTEXQVGIYIILSSSGDGAXXRDSGS 120
    |||||
Db 61 VSSSCGPOCHKGTPPTYKEAKQYLSYETLYANGSRTEXQVGIYIILSSSGDGAXXRDSGS 120
    |||||

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLTGCTGTLVAEXHVLTAACHIDG 180
    |||||
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLTGCTGTLVAEXHVLTAACHIDG 180
    |||||

QY 181 KTYVKGTKLRVGFLLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240
    |||||
Db 181 KTYVKGTKLRVGFLLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240
    |||||

QY 241 IGMDDYALLELKKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240
    |||||
Db 241 IGMDDYALLELKKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240
    |||||

QY 301 TYDLYQQCDAQPGASGYGVYVVMKRWQKQKWERKIIGIFSGHWDMMNGSPQDFNVAVR 360
    |||||
Db 301 TYDLYQQCDAQPGASGYGVYVVMKRWQKQKWERKIIGIFSGHWDMMNGSPQDFNVAVR 360
    |||||

QY 361 ITPLKYAQICYWIKGNLYDCREGDTVFLPGSN 392
    |||||
Db 361 ITPLKYAQICYWIKGNLYDCREGDTVFLPGSN 392
    |||||

RESULT 2
US-09-072-384-15
; Sequence 15, Application US/09072384
; Patent No. 6153420
; GENERAL INFORMATION:
; APPLICANT: Sheppard, Paul O.
; TITLE OF INVENTION: SERINE PROTEASE POLYPEPTIDES
; TITLE OF INVENTION: AND MATERIALS AND METHODS FOR MAKING THEM
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: ZymoGenetics, Inc.
; STREET: 1201 Eastlake Avenue East
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98102
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/072,384
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Parker, Gary E
; REGISTRATION NUMBER: 31,648
; REFERENCE/DOCKET NUMBER: 97-16C1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-442-6673
; TELEFAX: 206-442-6678
; TELEX:
; INFORMATION FOR SEQ ID NO: 15:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 392 amino acids
; TYPE: amino acid
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STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
FRAGMENT TYPE: internal
FEATURE:
NAME/KEY: Signal Sequence
LOCATION: 1...19
OTHER INFORMATION:
US-09-072-384-15

Query Match          98.7%; Score 2085; DB 3; Length 392;
Best Local Similarity 98.2%; Pred. No. 5.9e-216;
Matches 385; Conservative 1; Mismatches 6; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
    |||||
Db 1 MAGIPGLLFLFLCAVGVSPYSAPWKPPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
    |||||

QY 61 VSSSCGPOCHKGTPPTYKEAKQYLSYETLYANGSRTEXQVGIYIILSSSGDGAXXRDSGS 120
    |||||
Db 61 VSSSCGPOCHKGTPPTYKEAKQYLSYETLYANGSRTEXQVGIYIILSSSGDGAXXRDSGS 120
    |||||

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLTGCTGTLVAEXHVLTAACHIDG 180
    |||||
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLTGCTGTLVAEXHVLTAACHIDG 180
    |||||

QY 181 KTYVKGTKLRVGFLLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240
    |||||
Db 181 KTYVKGTKLRVGFLLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240
    |||||

QY 241 IGMDDYALLELKKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240
    |||||
Db 241 IGMDDYALLELKKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKNAND 240
    |||||

QY 301 TYDLYQQCDAQPGASGYGVYVVMKRWQKQKWERKIIGIFSGHWDMMNGSPQDFNVAVR 360
    |||||
Db 301 TYDLYQQCDAQPGASGYGVYVVMKRWQKQKWERKIIGIFSGHWDMMNGSPQDFNVAVR 360
    |||||

QY 361 ITPLKYAQICYWIKGNLYDCREGDTVFLPGSN 392
    |||||
Db 361 ITPLKYAQICYWIKGNLYDCREGDTVFLPGSN 392
    |||||

RESULT 3
US-09-072-384-18
; Sequence 18, Application US/09072384
; Patent No. 6153420
; GENERAL INFORMATION:
; APPLICANT: Sheppard, Paul O.
; TITLE OF INVENTION: SERINE PROTEASE POLYPEPTIDES
; TITLE OF INVENTION: AND MATERIALS AND METHODS FOR MAKING THEM
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: ZymoGenetics, Inc.
; STREET: 1201 Eastlake Avenue East
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98102
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/072,384
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Parker, Gary E
```


REGISTRATION NUMBER: 31,648
REFERENCE/DOCKET NUMBER: 97-16C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-442-6673
TELEFAX: 206-442-6678
TELEX:

INFORMATION FOR SEQ ID NO: 18:

SEQUENCE CHARACTERISTICS:

LENGTH: 383 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: protein

FRAGMENT TYPE: internal

FEATURE:

NAME/KEY: Signal Sequence

LOCATION: 1...19

OTHER INFORMATION:

US-09-072-384-18

Query Match 96.8%; Score 2044; DB 3; Length 383;
Best Local Similarity 98.4%; Pred. No. 1.5e-211;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCVAVGVSPYSAPWKTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLLCVAVGVSPYSAPWKTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPOCHKGTPLTYKEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXXRDSGS 120
DB 61 VSSSCGPOCHKGTPLTYKEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXXRDSGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLTVAEXHVLTAACHIDG 180
DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLTVAEXHVLTAACHIDG 180

QY 181 KTYVKGTKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240
DB 181 KTYVKGTKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240

QY 241 IGMDDYVALLLELKKPKRKFEMKICVSPAKQLPGRIHFSGYNDNRGNLVYRFDVKDE 300
DB 241 IGMDDYVALLLELKKPKRKFEMKICVSPAKQLPGRIHFSGYNDNRGNLVYRFDVKDE 300

QY 301 TYDLLYQCCDAQPGASGYVYVWKRQKQKWERKIIGFSGHQWDMNGSPQDFNVAVR 360
DB 301 TYDLLYQCCDAQPGASGYVYVWKRQKQKWERKIIGFSGHQWDMNGSPQDFNVAVR 360

QY 361 ITPLKVAQICYWKIGNYLDREG 383
DB 361 ITPLKVAQICYWKIGNYLDREG 383

RESULT 4

US-09-907-794A-261
Sequence 261, Application US/09907794A

Patent No. 6635468

GENERAL INFORMATION:

APPLICANT: Genentech, Inc.

APPLICANT: Ashkenazi, Avi

APPLICANT: Botstein, David

APPLICANT: Desnoyers, Luc

APPLICANT: Eaton, Dan L.

APPLICANT: Ferrara, Napoleone

APPLICANT: Filvaroff, Ellen

APPLICANT: Fong, Sherman

APPLICANT: Gao, Wei-Qiang

APPLICANT: Gerber, Hanspeter

APPLICANT: Grittisen, Mary E.

APPLICANT: Goddard, A.

APPLICANT: Godowski, Paul J.

APPLICANT: Grimaldi, Christopher J.

APPLICANT: Gurney, Austin L.

APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/907,794A
CURRENT FILING DATE: 2001-07-17
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/30095
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-907-794A-261

Query Match 96.8%; Score 2044; DB 4; Length 383;
Best Local Similarity 98.4%; Pred. No. 1.5e-211;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCVAVGVSPYSAPWKTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLLCVAVGVSPYSAPWKTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPOCHKGTPLTYKEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXXRDSGS 120
DB 61 VSSSCGPOCHKGTPLTYKEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXXRDSGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLTVAEXHVLTAACHIDG 180
DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLTVAEXHVLTAACHIDG 180

QY 181 KTYVKGTKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240

Db 181 KTVVGTQKLRVGLFKPFGXGGRGANDSTSAPEQKQFQWIRVXRTHVPKGIKGNAND 240
QY 241 IGMDDYDYLLELKKPKRKFPMKIGVSPPAKQLPGRRIHFSGYDNDPRGNLVYRFCVDKDE 300
Db 241 IGMDDYDYLLELKKPKRKFPMKIGVSPPAKQLPGRRIHFSGYDNDPRGNLVYRFCVDKDE 300
QY 301 TYDLLYQCCDAQPGASGVYVYRMWKRQOQKWERKIIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAQPGASGVYVYRMWKRQOQKWERKIIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPKYAQICYWIKNYLDREG 383
Db 361 ITPKYAQICYWIKNYLDREG 383

RESULT 5
US-09-905-125A-261
; Sequence 261, Application US/09905125A
; Patent No. 6664376
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,125A
; PRIOR FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30

; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-905-125A-261
Query Match 96.8%; Score 2044; DB 4; Length 383;
Best Local Similarity 98.4%; Pred. No. 1.5e-211;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;
QY 1 MAGIPGLLELLEFLICAVQVSPYSAPWKPTWPAYRLPVVLPOSTLNIAKPDGAEAKLE 60
Db 1 MAGIPGLLELLEFLICAVQVSPYSAPWKPTWPAYRLPVVLPOSTLNIAKPDGAEAKLE 60
QY 61 VSSSCGPOCHKGTPLPTYKEAKQYLSYETLYANGSRTEXQVGIYILSSSGDGAXXRDSGS 120
Db 61 VSSSCGPOCHKGTPLPTYEEAKQYLSYETLYANGSRTEQVGIYILSSSGDGQAHRDSSG 120
QY 121 SGKSRKQIYGYDSRFSIFGKDFLLNYPFSTS VKLSTGCTGTLVAEXHVLTAACHIDG 180
Db 121 SGKSRKQIYGYDSRFSIFGKDFLLNYPFSTS VKLSTGCTGTLVAEXHVLTAACHIDG 180
QY 181 KTYVGTQKLRVGLFKPFGXGGRGANDSTSAPEQKQFQWIRVXRTHVPKGIKGNAND 240
Db 181 KTYVGTQKLRVGLFKPFGXGGRGANDSTSAPEQKQFQWIRVXRTHVPKGIKGNAND 240
QY 241 IGMDDYDYLLELKKPKRKFPMKIGVSPPAKQLPGRRIHFSGYDNDPRGNLVYRFCVDKDE 300
Db 241 IGMDDYDYLLELKKPKRKFPMKIGVSPPAKQLPGRRIHFSGYDNDPRGNLVYRFCVDKDE 300
QY 301 TYDLLYQCCDAQPGASGVYVYRMWKRQOQKWERKIIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAQPGASGVYVYRMWKRQOQKWERKIIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPKYAQICYWIKNYLDREG 383
Db 361 ITPKYAQICYWIKNYLDREG 383
RESULT 6
US-09-902-775A-261
; Sequence 261, Application US/09902775A
; Patent No. 6686451
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.

APPLICANT: Kijavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT FILING DATE: 2001-07-10
CURRENT APPLICATION NUMBER: US/09/902.775A
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/30095
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-902-775A-261

Query Match 96.8%; Score 2044; DB 4; Length 383;
Best Local Similarity 98.4%; Pred. No. 1.5e-211;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGTGLLLFLLLCVAGGVSPYSAPWKPETWPAYRLPVVLPOSTLNLAKEPFGAEAKLE 60
DB 1 MAGTGLLLFLLLCVAGGVSPYSAPWKPETWPAYRLPVVLPOSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPOCHKGTPLPTYEAKQVLSYETLYANGSRTEKQVGIYILSSGDXRDSGS 120
DB 61 VSSSCGPOCHKGTPLPTYEAKQVLSYETLYANGSRTEKQVGIYILSSGDXRDSGS 120
QY 121 SGKSRKRIQYIDSRFSIFGKDFLLNYPFSTSVKLSGTGTLVAEXHVLTAACHIDG 180
DB 121 SGKSRKRIQYIDSRFSIFGKDFLLNYPFSTSVKLSGTGTLVAEXHVLTAACHIDG 180
QY 181 KTYVGTQKLRVGLPKPKFQKGGGRANDSTSAMPEQMKFQWIRVKTHTVPGKWIKNAND 240

DB 181 KTYVGTQKLRVGLPKPKFQKGGGRANDSTSAMPEQMKFQWIRVKTHTVPGKWIKNAND 240
QY 241 IGMDDYVALLLELKKPKHKKFKMGKIGVSPPAKOLPGGRIFHFGYDNDRCNLVYRCDVKDE 300
DB 241 IGMDDYVALLLELKKPKHKKFKMGKIGVSPPAKOLPGGRIFHFGYDNDRCNLVYRCDVKDE 300
QY 301 TYDLLYQOCDAQPGASGYGVYVVMWKRQOQKWKRIIGIFSGHWDVMDGSPQDFNVAVR 360
DB 301 TYDLLYQOCDAQPGASGYGVYVVMWKRQOQKWKRIIGIFSGHWDVMDGSPQDFNVAVR 360
QY 361 ITELKVAQICWIKGNLYDCREG 383
DB 361 ITELKVAQICWIKGNLYDCREG 383

RESULT 7
US-09-906-700-261
Sequence 261, Application US/09906700
Patent No. 6723535
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kijavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/906,700
CURRENT FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564

; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-906-700-261

Query Match 96.8%; Score 2044; DB 4; Length 383;
Best Local Similarity 98.4%; Pred. No. 1.5e-211;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAGGVSPYSAPWKPPTWPAVRLPVVLPQSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLLCAGGVSPYSAPWKPPTWPAVRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPGQCHKGTPLPTTYEAKQYLSYETLYANGSRTEXQVGIYIILSSSGDGAXXRDSGS 120
DB 61 VSSSCGPGQCHKGTPLPTTYEAKQYLSYETLYANGSRTEXQVGIYIILSSSGDGAXXRDSGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEXHVLTAACHIDG 180
DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEXHVLTAACHIDG 180

QY 181 KTYVKGTKLRVGLFKPKDGGGRANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240
DB 181 KTYVKGTKLRVGLFKPKDGGGRANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240

QY 241 IGMDDYALLESKPKHFKFMKIGVSPPAKQLPGGRHFGSGYDNDPRGNLVRFCVDKDE 300
DB 241 IGMDDYALLESKPKHFKFMKIGVSPPAKQLPGGRHFGSGYDNDPRGNLVRFCVDKDE 300

QY 301 TYDLLYQQCDAQSGAGSYGVYVMWRKQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
DB 301 TYDLLYQQCDAQSGAGSYGVYVMWRKQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360

QY 361 ITPKYAQICYWIKGNLYDCREG 383
DB 361 ITPKYAQICYWIKGNLYDCREG 383

RESULT 8
US-09-903-603A-261
; Sequence 261, Application US/09903603A
; Patent No. 6767995
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.

; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: GNE.1618P2C12
; CURRENT APPLICATION NUMBER: US/09/903,603A
; CURRENT FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-903-603A-261

Query Match 96.8%; Score 2044; DB 4; Length 383;
Best Local Similarity 98.4%; Pred. No. 1.5e-211;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAGGVSPYSAPWKPPTWPAVRLPVVLPQSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLLCAGGVSPYSAPWKPPTWPAVRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPGQCHKGTPLPTTYEAKQYLSYETLYANGSRTEXQVGIYIILSSSGDGAXXRDSGS 120
DB 61 VSSSCGPGQCHKGTPLPTTYEAKQYLSYETLYANGSRTEXQVGIYIILSSSGDGAXXRDSGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEXHVLTAACHIDG 180
DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEXHVLTAACHIDG 180

QY 181 KTYVKGTKLRVGLFKPKDGGGRANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240
DB 181 KTYVKGTKLRVGLFKPKDGGGRANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240

QY 241 IGMDYDVALLELKKPHKFKFKVSPKQALPGCRHFGSGYDNDRGNLVYRFDVKDE 300
 Db 241 IGMDYDVALLELKKPHKFKFKVSPKQALPGCRHFGSGYDNDRGNLVYRFDVKDE 300
 QY 301 TYDLYQQCDAOPGASGYVYRWMKQKQKWERKIIGIFSGHQWVDMGSPQDFNVAVR 360
 Db 301 TYDLYQQCDAOPGASGYVYRWMKQKQKWERKIIGIFSGHQWVDMGSPQDFNVAVR 360
 QY 361 ITPLKYAQICYWIKGNLYDCREG 383
 Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 9

US-09-904-920A-261
 ; Sequence 261, Application US/09904920A
 ; Patent No. 6806352
 ; GENERAL INFORMATION:
 ; APPLICANT: Genentech, Inc.
 ; APPLICANT: Ashkenazi, Avi
 ; APPLICANT: Botstein, David
 ; APPLICANT: Desnoyers, Luc
 ; APPLICANT: Eaton, Dan L.
 ; APPLICANT: Ferrara, Napoleone
 ; APPLICANT: Filvaroff, Ellen
 ; APPLICANT: Fong, Sherman
 ; APPLICANT: Gao, Wei-Qiang
 ; APPLICANT: Gerber, Hanspeter
 ; APPLICANT: Gerritsen, Mary E.
 ; APPLICANT: Goddard, A.
 ; APPLICANT: Godowski, Paul J.
 ; APPLICANT: Grimaldi, Christopher J.
 ; APPLICANT: Gurney, Austin L.
 ; APPLICANT: Hillan, Kenneth, J.
 ; APPLICANT: Kljavin, Ivar J.
 ; APPLICANT: Mather, Jennie P.
 ; APPLICANT: Pan, James
 ; APPLICANT: Paoni, Nicholas F.
 ; APPLICANT: Roy, Margaret Ann
 ; APPLICANT: Stewart, Timothy A.
 ; APPLICANT: Tumas, Daniel
 ; APPLICANT: Williams, P. Mickey
 ; APPLICANT: Wood, William, I.
 ; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
 ; FILE OF INVENTION: Acids Encoding the Same
 ; FILE REFERENCE: 10466-14
 ; CURRENT APPLICATION NUMBER: US/09/904,920A
 ; CURRENT FILING DATE: 2001-07-13
 ; PRIOR APPLICATION NUMBER: PCT/US00/04414
 ; PRIOR FILING DATE: 2000-02-22
 ; PRIOR APPLICATION NUMBER: US 60/143,048
 ; PRIOR FILING DATE: 1999-07-07
 ; PRIOR APPLICATION NUMBER: US 60/145,698
 ; PRIOR FILING DATE: 1999-07-26
 ; PRIOR APPLICATION NUMBER: US 60/146,222
 ; PRIOR FILING DATE: 1999-07-28
 ; PRIOR APPLICATION NUMBER: PCT/US99/20594
 ; PRIOR FILING DATE: 1999-09-08
 ; PRIOR APPLICATION NUMBER: PCT/US99/20944
 ; PRIOR FILING DATE: 1999-09-13
 ; PRIOR APPLICATION NUMBER: PCT/US99/21090
 ; PRIOR FILING DATE: 1999-09-15
 ; PRIOR APPLICATION NUMBER: PCT/US99/21547
 ; PRIOR FILING DATE: 1999-09-15
 ; PRIOR APPLICATION NUMBER: PCT/US99/23089
 ; PRIOR FILING DATE: 1999-10-05
 ; PRIOR APPLICATION NUMBER: PCT/US99/28214
 ; PRIOR FILING DATE: 1999-11-29
 ; PRIOR APPLICATION NUMBER: PCT/US99/28313
 ; PRIOR FILING DATE: 1999-11-30
 ; PRIOR APPLICATION NUMBER: PCT/US99/28564
 ; PRIOR FILING DATE: 1999-12-02

; PRIOR APPLICATION NUMBER: PCT/US99/28565
 ; PRIOR FILING DATE: 1999-12-02
 ; PRIOR APPLICATION NUMBER: PCT/US99/30095
 ; PRIOR FILING DATE: 1999-12-16
 ; PRIOR APPLICATION NUMBER: PCT/US99/30911
 ; PRIOR FILING DATE: 1999-12-20
 ; PRIOR APPLICATION NUMBER: PCT/US99/30999
 ; PRIOR FILING DATE: 1999-12-20
 ; PRIOR APPLICATION NUMBER: PCT/US00/00219
 ; PRIOR FILING DATE: 2000-01-05
 ; NUMBER OF SEQ ID NOS: 423
 ; SEQ ID NO 261
 ; LENGTH: 383
 ; TYPE: PRT
 ; ORGANISM: Homo Sapien
 US-09-904-920A-261

Query Match 96.8%; Score 2044; DB 4; Length 383;

Best Local Similarity 98.4%; Pred. No. 1.5e-211;

Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFPLLCAVGQVSPYSAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
 Db 1 MAGIPGLLFLFPLLCAVGQVSPYSAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
 QY 61 VSSCGPQCHKGTPPTYKEAQYLSYETLYANGSRTEQVGIYIILSSGCGAXXRDSGS 120
 Db 61 VSSCGPQCHKGTPPTYBEAQYLSYETLYANGSRTEQVGIYIILSSGCGAXXRDSGS 120
 QY 121 SGKSRKROIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
 Db 121 SGKSRKROIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
 QY 181 KTYVKTQKLRVGFLLKPKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
 Db 181 KTYVKTQKLRVGFLLKPKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
 QY 241 IGMDYDVALLELKKPHKFKFKVSPKQALPGCRHFGSGYDNDRGNLVYRFDVKDE 300
 Db 241 IGMDYDVALLELKKPHKFKFKVSPKQALPGCRHFGSGYDNDRGNLVYRFDVKDE 300
 QY 301 TYDLYQQCDAOPGASGYVYRWMKQKQKWERKIIGIFSGHQWVDMGSPQDFNVAVR 360
 Db 301 TYDLYQQCDAOPGASGYVYRWMKQKQKWERKIIGIFSGHQWVDMGSPQDFNVAVR 360
 QY 361 ITPLKYAQICYWIKGNLYDCREG 383
 Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 10

US-09-909-064-261
 ; Sequence 261, Application US/09909064
 ; Patent No. 6818449
 ; GENERAL INFORMATION:
 ; APPLICANT: Genentech, Inc.
 ; APPLICANT: Ashkenazi, Avi
 ; APPLICANT: Botstein, David
 ; APPLICANT: Desnoyers, Luc
 ; APPLICANT: Eaton, Dan L.
 ; APPLICANT: Ferrara, Napoleone
 ; APPLICANT: Filvaroff, Ellen
 ; APPLICANT: Fong, Sherman
 ; APPLICANT: Gao, Wei-Qiang
 ; APPLICANT: Gerber, Hanspeter
 ; APPLICANT: Gerritsen, Mary E.
 ; APPLICANT: Goddard, A.
 ; APPLICANT: Godowski, Paul J.
 ; APPLICANT: Grimaldi, Christopher J.
 ; APPLICANT: Gurney, Austin L.
 ; APPLICANT: Hillan, Kenneth, J.
 ; APPLICANT: Kljavin, Ivar J.
 ; APPLICANT: Mather, Jennie P.

APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/909,064
CURRENT FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/30095
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-909-064-261

Query Match 96.8%; Score 2044; DB 4; Length 383;
Best Local Similarity 98.4%; Pred. No. 1.5e-211;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFLLCAGGVSPVSPKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAGGVSPVSPKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

Qy 61 VSSSCGPGCHKGTPLPYKEAKQYLSYETLLYANGSRTEXQVGYIYILSSGSGDGAAXRDSGS 120
Db 61 VSSSCGPGCHKGTPLPYKEAKQYLSYETLLYANGSRTEXQVGYIYILSSGSGDGAAXRDSGS 120

Qy 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSGTGTLVAEXHVLTAACHIDG 180
Db 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSGTGTLVAEXHVLTAACHIDG 180

Qy 181 KTVYKGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
Db 181 KTVYKGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240

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Db 241 IGMVDYDYLLELKKPKRKFPMKIGVSPPAKQIPLGGRIHPSGYDNDPRLPGNLVYRFCVDKDE 300

Qy 301 TYDLLYQQCDAOPGASGVGVYVMMKROQKQKWERKIIGIFSGHGWYDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAOPGASGVGVYVMMKROQKQKWERKIIGIFSGHGWYDMNGSPQDFNVAVR 360

Qy 361 ITPLKYAQICYWIKGNLYLDCREG 383
Db 361 ITPLKYAQICYWIKGNLYLDCREG 383

RESULT 11
US-09-905-381A-261
Sequence 261, Application US/09905381A
Patent No. 6818746
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/905,381A
CURRENT FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565

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; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-905-381A-261

Query Match          96.8%; Score 2044; DB 4; Length 383;
Best Local Similarity 98.4%; Pred. No. 1.5e-211;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPQCHKGTPPTYKEAKQVLSYETLYANGSRTEXOVGIYIILSSGDXGAXRDSGS 120
Db 61 VSSSCGPQCHKGTPPTYKEAKQVLSYETLYANGSRTEXOVGIYIILSSGDXGAXRDSGS 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEXHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEXHVLTAACHIDG 180
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Db 181 KTYVKGTKLRVGFLLKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240
QY 241 IGMVDYVALLLELKKPHKRFKMGKIGVSPAPKQLPGGRIFHSGYDNDPRGNLVYRFDVKDE 300
Db 241 IGMVDYVALLLELKKPHKRFKMGKIGVSPAPKQLPGGRIFHSGYDNDPRGNLVYRFDVKDE 300
QY 301 TYDLLYQQCDAQPGASGYGVYVWMKRRQKQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAQPGASGYGVYVWMKRRQKQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPKVAQICYWKGNVLDREG 383
Db 361 ITPKVAQICYWKGNVLDREG 383
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RESULT 12

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US-09-906-618-261
; Sequence 261, Application US/09906618
; Patent No. 6828146
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gertsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
```

```
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/906,618
; CURRENT FILING DATE: 2001-07-16
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-906-618-261

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Best Local Similarity 98.4%; Pred. No. 1.5e-211;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

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Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEXHVLTAACHIDG 180
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; CURRENT APPLICATION NUMBER: US/09/551,826D
; CURRENT FILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 316
; TYPE: PR1
; ORGANISM: Bacillus licheniformis
US-09-551-826D-2

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QY 120 SSGKSRK-----ROIYGYDSRFSIFGKDFLLNYPFSTSVKLST---GCTGLVAE 167
Db 77 TKEAEKSPAKAPYSIKSVIGSDRTRVTN---TTAPYRAIVHISSIGSCTGMMIGP 133

QY 168 XHVLTAACHIDGKT-YVKGTKLKVGLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKR 226
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QY 227 THVPKGIKGNANDIGMDYDIALLEKPKHKFKMKGIVSPPAKQLPGGRIHFSGYDNR 286
Db 176 YFIPSGWRSGNTN---YDYGAIELSEPIGNTVGFGYSYTTSLVGTGTTVTISGYPGDK 230

QY 287 PGNLVYRFGD--VKDETDLLYQQCDAPGASGYGVYVVMKRWKQOQKWERKII---GIF 340
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QY 341 SGHQQWDMNGSPQDFNVAVRITPLKYAICYW 372
Db 290 GG-----SSYNRGTRITKEVDFDNLTNW 311

RESULT 15
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; Sequence 10, Application US/09551826D
; Patent No. 6558939
; GENERAL INFORMATION:
; APPLICANT: No. 6558939regaard-Madsen, Mads
; APPLICANT: Ostergaard, Peter Rahbek
; APPLICANT: Christensen, Claus Bo Voge
; APPLICANT: Lassen, Soren Flensted
; TITLE OF INVENTION: No. 6558939el Proteases And Variants Thereof
; FILE REFERENCE: 5665.200-US
; CURRENT APPLICATION NUMBER: US/09/551,826D
; CURRENT FILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 10
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; TYPE: PR1
; ORGANISM: Bacillus licheniformis CDJ31
US-09-551-826D-10

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; CURRENT APPLICATION NUMBER: US/09/551,826D
; CURRENT FILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 14
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; LENGTH: 314
; TYPE: PR1
; ORGANISM: Bacillus licheniformis AC116
US-09-551-826D-6

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Best Local Similarity 24.4%; Pred. No. 8.9e-12;
Matches 78; Conservative 36; Mismatches 141; Indels 65; Gaps 14;

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QY 123 KSRKRIQYGYDSRFSIFGKDFLLNYPFSTSVKLST---GCTGLVAEXHVLTAACHID 179
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QY 180 -GKTYVKGTKLKVGLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNGA 238
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QY 297 VKDETDLLYQQCDAPGASGYGVYVVMKRWKQOQKWERKII---GIFSGHQWDMNGSP 352
Db 241 AVSETYKLOY-AIDTYGGQSGSPVFEKSSRTNCSGPCSLAVHTNGVYGG----- 289

QY 353 QDFNVAVRITPLKYAICYW 372
Db 290 SSYNRGTRITKEVDFDNLTSW 309

RESULT 14
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; Sequence 2, Application US/09551826D
; Patent No. 6558939
; GENERAL INFORMATION:
; APPLICANT: No. 6558939regaard-Madsen, Mads
; APPLICANT: Ostergaard, Peter Rahbek
; APPLICANT: Christensen, Claus Bo Voge
; APPLICANT: Lassen, Soren Flensted
; TITLE OF INVENTION: No. 6558939el Proteases And Variants Thereof
; FILE REFERENCE: 5665.200-US
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Db	242	GNIAV-----SETYKLOY-AIDTYGGSGSPVYEASSRTNCSGPCSLAVHTNGVYGG-	293
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: July 1, 2005, 21:08:25 ; Search time 90.3582 Seconds
(without alignments)
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Title: US-09-658-677-2
Perfect score: 2112
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Searched: 1722976 seqs, 385795295 residues

Total number of hits satisfying chosen parameters: 1722976

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
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Listing first 45 summaries

Database : Published Applications AA:
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22: /cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pcp:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2	2044	96.8	383	9	US-09-309-320-261
3	2044	96.8	383	9	US-09-909-088B-261
4	2044	96.8	383	9	US-09-905-291A-261
5	2044	96.8	383	9	US-09-902-853-261
6	2044	96.8	383	9	US-09-907-824-261
7	2044	96.8	383	9	US-09-907-841-261
8	2044	96.8	383	10	US-09-904-011-261
9	2044	96.8	383	10	US-09-903-640-261
10	2044	96.8	383	10	US-09-908-093-261
11	2044	96.8	383	10	US-09-906-742-261

12	2044	96.8	383	10	US-09-906-838-261	Sequence 261, App
13	2044	96.8	383	10	US-09-907-613-261	Sequence 261, App
14	2044	96.8	383	10	US-09-907-942-261	Sequence 261, App
15	2044	96.8	383	10	US-09-904-859-261	Sequence 261, App
16	2044	96.8	383	10	US-09-909-204-261	Sequence 261, App
17	2044	96.8	383	10	US-09-904-820-261	Sequence 261, App
18	2044	96.8	383	10	US-09-904-786-261	Sequence 261, App
19	2044	96.8	383	10	US-09-906-646-261	Sequence 261, App
20	2044	96.8	383	10	US-09-906-700-261	Sequence 261, App
21	2044	96.8	383	10	US-09-903-786-261	Sequence 261, App
22	2044	96.8	383	10	US-09-902-903-261	Sequence 261, App
23	2044	96.8	383	10	US-09-903-749A-261	Sequence 261, App
24	2044	96.8	383	10	US-09-904-956-261	Sequence 261, App
25	2044	96.8	383	10	US-09-902-736-261	Sequence 261, App
26	2044	96.8	383	10	US-09-907-794-261	Sequence 261, App
27	2044	96.8	383	10	US-09-903-943-261	Sequence 261, App
28	2044	96.8	383	10	US-09-907-925-261	Sequence 261, App
29	2044	96.8	383	10	US-09-904-462-261	Sequence 261, App
30	2044	96.8	383	10	US-09-902-692-261	Sequence 261, App
31	2044	96.8	383	10	US-09-903-520-261	Sequence 261, App
32	2044	96.8	383	10	US-09-903-056-261	Sequence 261, App
33	2044	96.8	383	10	US-09-984-130-45	Sequence 45, App1
34	2044	96.8	383	10	US-09-984-130-126	Sequence 126, App
35	2044	96.8	383	10	US-09-909-064-261	Sequence 261, App
36	2044	96.8	383	10	US-09-904-553-261	Sequence 261, App
37	2044	96.8	383	10	US-09-905-381-261	Sequence 261, App
38	2044	96.8	383	10	US-09-904-485-261	Sequence 261, App
39	2044	96.8	383	10	US-09-905-348-261	Sequence 261, App
40	2044	96.8	383	10	US-09-905-088-261	Sequence 261, App
41	2044	96.8	383	10	US-09-907-575-261	Sequence 261, App
42	2044	96.8	383	10	US-09-905-075-261	Sequence 261, App
43	2044	96.8	383	10	US-09-902-759-261	Sequence 261, App
44	2044	96.8	383	10	US-09-902-634-261	Sequence 261, App
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ALIGNMENTS

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; Sequence 12, Application US/09765205
; Patent No. US20020034800A1
; GENERAL INFORMATION:
; APPLICANT: Cao, Li
; TITLE OF INVENTION: BONE MARROW SECRETED PROTEINS AND POLYNUCLEOTIDES
; FILE REFERENCE: 1458.004/200130.449
; CURRENT APPLICATION NUMBER: US/09/765,205
; CURRENT FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: US/09/212,440
; PRIOR FILING DATE: 1998-12-16
; NUMBER OF SEQ ID NOS: 46
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 12
; LENGTH: 383
; TYPE: PRT
; ORGANISM: human
; US-09-765-205-12

Query Match	96.8%	Score 2044;	DB 9;	Length 383;
Best Local Similarity	98.4%;	Pred. No. 4e-193;		
Matches 377;	Conservative 1;	Mismatches 5;	Indels 0;	Gaps 0;
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Db	61	VSSSCGPQCHKGTPL	PTYEAKQYLSYETLYANGSTEXQVGIYILSSSGDGA	XXRDSGS 120
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RESULT 2
US-09-909-320-261
; Sequence 261, Application US/09909320
; Patent No. US20020132240A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/909,320
; CURRENT FILING DATE: 2002-01-04
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
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; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
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; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
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; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
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; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-909-320-261
Query Match 96.8%; Score 2044; DB 9; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193; 5; Indels 0; Gaps 0;
Matches 377; Conservative 1; Mismatches 5;
Qy 1 MAGIPLGLFLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLQSTLNLAQPDGAEAKLE 60
Db 1 MAGIPLGLFLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLQSTLNLAQPDGAEAKLE 60
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; Sequence 261, Application US/09909088B
; Patent No. US20020146709A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.

```

; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/909/088B
; PRIOR FILING DATE: 2001-07-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-909-088B-261

Query Match          96.8%; Score 2044; DB 9; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFLLCVAGQVSPYAPKPTWPAAYRLPVVLPQSTLNLAKEPFGABAKLE 60
Db 1 MAGIPGLLFLFLLCVAGQVSPYAPKPTWPAAYRLPVVLPQSTLNLAKEPFGABAKLE 60
Qy 61 VSSSCGQCHGTPLPTPYEAKQVLSYETLYANGSRTEQVGIYILSSSGDGAXRDSGS 120
Db 61 VSSSCGQCHGTPLPTPYEAKQVLSYETLYANGSRTEQVGIYILSSSGDGAXRDSGS 120
Qy 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSCTGCTTLVAEKHVLTAACHIDHG 180
Db 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSCTGCTTLVAEKHVLTAACHIDHG 180

; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,291A
; PRIOR FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
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; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-905-291A-261

; Sequence 261, Application US/09905291A
; Patent No. US20020160374A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,291A
; PRIOR FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
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; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
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; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-905-291A-261

Query Match 96.8%; Score 2044; DB 9; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193; 5; Indels 0; Gaps 0;
Matches 377; Conservative 1; Mismatches 0;

QY 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSTEXQVGIYILSSSGDGAHXRDGSG 120
Db 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSTEXQVGIYILSSSGDGAHXRDGSG 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLNYPFSTSVKLSCTGTLVAEXHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLNYPFSTSVKLSCTGTLVAEXHVLTAACHIDG 180
QY 181 KTVVKGTKLRVGLFKPKFKDGGRGANDSTSAPEQMKFQIRVKRTHVPKGIKNAND 240
Db 181 KTVVKGTKLRVGLFKPKFKDGGRGANDSTSAPEQMKFQIRVKRTHVPKGIKNAND 240
QY 241 IGMVDYALLEKKPKHKPKMKIGVSPPAKQLPGRIHFSGYNDPRGNLVYRFDVKDE 300
Db 241 IGMVDYALLEKKPKHKPKMKIGVSPPAKQLPGRIHFSGYNDPRGNLVYRFDVKDE 300
QY 301 TYDLLYQCCDAQPGASGYVYVWRKQKQKWKRIIGIFSGHQWVDMNGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAQPGASGYVYVWRKQKQKWKRIIGIFSGHQWVDMNGSPQDFNVAVR 360
QY 361 ITPLKVAQICYKGNLYDCREG 383
Db 361 ITPLKVAQICYKGNLYDCREG 383

RESULT 5
US-09-902-853-261
; Publication No. US0902853
; Publication No. US20020192659A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.

; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/902,853
; CURRENT FILING DATE: 2001-07-10
; PRIOR APPLICATION NUMBER: US/09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-902-853-261

Query Match 96.8%; Score 2044; DB 9; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193; 5; Indels 0; Gaps 0;
Matches 377; Conservative 1; Mismatches 0;

QY 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSTEXQVGIYILSSSGDGAHXRDGSG 120
Db 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSTEXQVGIYILSSSGDGAHXRDGSG 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLNYPFSTSVKLSCTGTLVAEXHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLNYPFSTSVKLSCTGTLVAEXHVLTAACHIDG 180

[illegible]

RESULT 6

RESULT 6

US-09-907-824-261

Sequence 261, Application US/09907824

Publication No. US20020197671A1

GENERAL INFORMATION:

APPLICANT: Genentech, Inc.

APPLICANT: Ashkenazi, Avi

APPLICANT: Botstein, David

APPLICANT: Desnoyers, Luc

APPLICANT: Eaton, Dan L.

APPLICANT: Ferrara, Napoleone

APPLICANT: Filvaroff, Ellen

APPLICANT: Fong, Sherman

APPLICANT: Gao, Wei-Qiang

APPLICANT: Gerber, Hanspeter

APPLICANT: Gerritsen, Mary E.

APPLICANT: Goddard, A.

APPLICANT: Godowski, Paul J.

APPLICANT: Grimaldi, Christopher J.

APPLICANT: Gurney, Austin L.

APPLICANT: Hillan, Kenneth, J.

APPLICANT: Kljavin, Ivar J.

APPLICANT: Mather, Jennie P.

APPLICANT: Pan, James

APPLICANT: Paoni, Nicholas F.

APPLICANT: Roy, Margaret Ann

APPLICANT: Stewart, Timothy A.

APPLICANT: Tumas, Daniel

APPLICANT: Williams, P. Mickey

APPLICANT: Wood, William, I.

TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic

TITLE OF INVENTION: Acids Encoding the Same

FILE REFERENCE: 10466-14

CURRENT APPLICATION NUMBER: US/09/907,824

CURRENT FILING DATE: 2001-07-17

PRIOR APPLICATION NUMBER: 09/665,350

PRIOR FILING DATE: 2000-09-18

PRIOR APPLICATION NUMBER: PCT/US00/04414

PRIOR FILING DATE: 2000-02-22

PRIOR APPLICATION NUMBER: US 60/143,048

PRIOR FILING DATE: 1999-07-07

PRIOR APPLICATION NUMBER: US 60/145,698

PRIOR FILING DATE: 1999-07-26

PRIOR APPLICATION NUMBER: US 60/146,222

PRIOR FILING DATE: 1999-07-28

PRIOR APPLICATION NUMBER: PCT/US99/20594

PRIOR FILING DATE: 1999-09-08

PRIOR APPLICATION NUMBER: PCT/US99/20944

PRIOR FILING DATE: 1999-09-13

PRIOR APPLICATION NUMBER: PCT/US99/21090

PRIOR FILING DATE: 1999-09-15

PRIOR APPLICATION NUMBER: PCT/US99/21547

PRIOR FILING DATE: 1999-09-15

PRIOR APPLICATION NUMBER: PCT/US99/23089

PRIOR FILING DATE: 1999-10-05

PRIOR APPLICATION NUMBER: PCT/US99/28214

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; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-907-824-261

Query Match          96.8%; Score 2044; DB 9; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0

Qy 1 MAGIPGLLFLLLFLLCAVGQVSPYSAPWKPTWPAYRLPVVLPOSTNLAKPDFGAEAKLE 60
Db 1 MAGIPGLLFLLLFLLCAVGQVSPYSAPWKPTWPAYRLPVVLPOSTNLAKPDFGAEAKLE 60

Qy 61 VSSSCGPQCHKGTPLPTYEAKQVLSYETLIYANGSRTEQVGIYILSSGDXRRDSGS 120
Db 61 VSSSCGPQCHKGTPLPTYEAKQVLSYETLIYANGSRTEQVGIYILSSGDXRRDSGS 120

Qy 121 SGKSRKRQIYGYDSRFSIIFGKDFLLNYPFSTSVKLSGTGCTGLVAEXHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIIFGKDFLLNYPFSTSVKLSGTGCTGLVAEXHVLTAACHIDG 180

Qy 181 KTVYKGTQKLRVGLPKPKFDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGMWIKGNAND 240
Db 181 KTVYKGTQKLRVGLPKPKFDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGMWIKGNAND 240

Qy 241 IGMDDYVALLLELKKPKRKFEMKIGVSPAPKQLPGRIHFSGYDNDRPGNLVYRFDVKDE 300
Db 241 IGMDDYVALLLELKKPKRKFEMKIGVSPAPKQLPGRIHFSGYDNDRPGNLVYRFDVKDE 300

Qy 301 TYDLLYQQCDAQPCAGSGYGYVVRWKRRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAQPCAGSGYGYVVRWKRRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360

Qy 361 ITPKVAQICYWIKNYLDRCRG 383
Db 361 ITPKVAQICYWIKNYLDRCRG 383

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RESIN, T 7

US-09-097-841-261
 ; Sequence 261, Application US/09907841
 ; Publication No. US20020198366A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Genentech, Inc.
 ; APPLICANT: Ashkenazi, Avi
 ; APPLICANT: Botstein, David
 ; APPLICANT: Desnovers, Luc
 ; APPLICANT: Eaton, Dan L.
 ; APPLICANT: Ferrara, Napoleone
 ; APPLICANT: Filvaroff, Ellen
 ; APPLICANT: Fong, Sherman
 ; APPLICANT: Gao, Wei-Qiang
 ; APPLICANT: Garber, Hanspeter
 ; APPLICANT: Gerritsen, Mary E.
 ; APPLICANT: Goddard, A.
 ; APPLICANT: Godowski, Paul J.

APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/907,841
PRIOR FILING DATE: 2001-11-20
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 393
TYPE: PRT
ORGANISM: Homo Sapien
US-09-907-841-261

Query Match 96.8%; Score 2044; DB 9; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

Qy 1 MAGIPGLFLLFLLCAVGQVSPYSAPWKTWPAYRLPVVLPQSTLNLAQDFGAEAKLE 60
Db 1 MAGIPGLFLLFLLCAVGQVSPYSAPWKTWPAYRLPVVLPQSTLNLAQDFGAEAKLE 60

Qy 61 VSSSCGPGCHGTPLPTYKEAKQVLSYETLYANGSRTEQVGIYILSSGSGAXRDSGS 120
Db 61 VSSSCGPGCHGTPLPTYKEAKQVLSYETLYANGSRTEQVGIYILSSGSGAXRDSGS 120

Qy 121 SGSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
Db 121 SGSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180

Qy 181 KTVVGTQKLRVGLFKPKDFGGRGANDSTSAMPEQMKFQIRVKRTHVPKGIKGNAND 240
Db 181 KTVVGTQKLRVGLFKPKDFGGRGANDSTSAMPEQMKFQIRVKRTHVPKGIKGNAND 240

Qy 241 IGMDDYVALLLEKKPHKPKMKTVGSPPAKQLPGGRIFHSGYNDPRGNLVYRFDVKDE 300
Db 241 IGMDDYVALLLEKKPHKPKMKTVGSPPAKQLPGGRIFHSGYNDPRGNLVYRFDVKDE 300

Qy 301 TYDLLYQCDAPQACSGVGVYVWVKRQKQKWERKLIIGFSGHQWDMNGSPQDENVAVR 360
Db 301 TYDLLYQCDAPQACSGVGVYVWVKRQKQKWERKLIIGFSGHQWDMNGSPQDENVAVR 360

Qy 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 8
US-09-904-011-261
Sequence 261, Application US/09904011
Publication No. US2003000350A1
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/904,011
CURRENT FILING DATE: 2001-07-11
PRIOR APPLICATION NUMBER: 09/665,350
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/30095
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999

; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00019
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-904-011-261

Query Match 96.8%; Score 2044; DB 10; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY	1	MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
DB	1	MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
QY	61	VSSSCGPGQCHKGTPPTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXHRDGS	120
DB	61	VSSSCGPGQCHKGTPPTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXHRDGS	120
QY	121	SGSRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLTGCTGLVAEXHVLTAACHIDG	180
DB	121	SGSRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLTGCTGLVAEXHVLTAACHIDG	180
QY	181	KTVYKGTQKLRVGLFKPKFQDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND	240
DB	181	KTVYKGTQKLRVGLFKPKFQDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND	240
QY	241	IGMDYDYLLELKKPKFKFVKIGVSPPAKQLPGRIHFSGYDNDPGLNLYVYRCDVKDE	300
DB	241	IGMDYDYLLELKKPKFKFVKIGVSPPAKQLPGRIHFSGYDNDPGLNLYVYRCDVKDE	300
QY	301	TYDLYQQCDAQPGASGYGVYVMWKRQOQKWERKIIGIFSGHQMVMGSPQDFNVAVR	360
DB	301	TYDLYQQCDAQPGASGYGVYVMWKRQOQKWERKIIGIFSGHQMVMGSPQDFNVAVR	360
QY	361	ITPLKYAQICYWIKGNLYDCREG	383
DB	361	ITPLKYAQICYWIKGNLYDCREG	383

RESULT 9

US-09-903-640-261
; Sequence 261, Application US/09903640
; Publication No. US20030017463A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.

; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/903,640
; PRIOR FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-903-640-261

Query Match 96.8%; Score 2044; DB 10; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY	1	MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
DB	1	MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
QY	61	VSSSCGPGQCHKGTPPTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXHRDGS	120
DB	61	VSSSCGPGQCHKGTPPTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAXHRDGS	120
QY	121	SGSRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLTGCTGLVAEXHVLTAACHIDG	180
DB	121	SGSRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLTGCTGLVAEXHVLTAACHIDG	180
QY	181	KTVYKGTQKLRVGLFKPKFQDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND	240
DB	181	KTVYKGTQKLRVGLFKPKFQDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND	240
QY	241	IGMDYDYLLELKKPKFKFVKIGVSPPAKQLPGRIHFSGYDNDPGLNLYVYRCDVKDE	300
DB	241	IGMDYDYLLELKKPKFKFVKIGVSPPAKQLPGRIHFSGYDNDPGLNLYVYRCDVKDE	300
QY	301	TYDLYQQCDAQPGASGYGVYVMWKRQOQKWERKIIGIFSGHQMVMGSPQDFNVAVR	360
DB	301	TYDLYQQCDAQPGASGYGVYVMWKRQOQKWERKIIGIFSGHQMVMGSPQDFNVAVR	360
QY	361	ITPLKYAQICYWIKGNLYDCREG	383
DB	361	ITPLKYAQICYWIKGNLYDCREG	383

RESULT 10

US-09-908-093-261
; Sequence 261, Application US/09908093
; Publication No. US20030017498A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann

```

; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/908,093
; CURRENT FILING DATE: 2001-07-17
; PRIOR FILING DATE: 2001-07-17
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-908-093-261

Query Match          96.8%; Score 2044; DB 10; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAGVGQSPVSPAPKPTWPAYRLPVVLPQSTLNLAKPDPGAEAKLE 60
DB 1 MAGIPGLLFLFLLCAGVGQSPVSPAPKPTWPAYRLPVVLPQSTLNLAKPDPGAEAKLE 60

QY 61 VSSSCGPOCHKGTPLPTYEAKQYLSYETLYANGSRTEXQVGYIYLLSSGSGDGAAXRDSGS 120
DB 61 VSSSCGPOCHKGTPLPTYEAKQYLSYETLYANGSRTEXQVGYIYLLSSGSGDGAAXRDSGS 120

QY 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLTGCTGLVAEXHVLTAACHIHOG 180
DB 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLTGCTGLVAEXHVLTAACHIHOG 180

QY 181 KTVVKGQKLRVGLFKPKDGGGRGNDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
DB 181 KTVVKGQKLRVGLFKPKDGGGRGNDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240

QY 241 IGMDDYDYLLELKKPKHKKFMKIGVSPPAKQLPGGRHIFSGYDNDRPGNLVYRFCVDKDE 300
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DB 241 IGMDDYDYLLELKKPKHKKFMKIGVSPPAKQLPGGRHIFSGYDNDRPGNLVYRFCVDKDE 300
QY 301 TYDLLYQQCDAOPGASGVYVYVMKRRQOQKWKRIIGIFSGHQWVDMNGSPQDFNVAVR 360
DB 301 TYDLLYQQCDAOPGASGVYVYVMKRRQOQKWKRIIGIFSGHQWVDMNGSPQDFNVAVR 360
QY 361 ITPKYAQICYWIKGNLYDCREG 383
DB 361 ITPKYAQICYWIKGNLYDCREG 383

RESULT 11
US-09-906-742-261
; Sequence 261, Application US/09906742
; Publication No. US20030023054A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/906,742
; CURRENT FILING DATE: 2001-07-16
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
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; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-906-742-261

Query Match          96.8%; Score 2044; DB 10; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPGCHKTGTPPTYKEAKOYLSYETLYANGSRTEXOVGIYILSSSGDGAXXRDSGS 120
Db 61 VSSSCGPGCHKTGTPPTYKEAKOYLSYETLYANGSRTEXOVGIYILSSSGDGAXXRDSGS 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
QY 181 KTVYGTQKLRVGLFKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTHTVPGKWIKNAND 240
Db 181 KTVYGTQKLRVGLFKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTHTVPGKWIKNAND 240
QY 241 IGMDDYVALLLEKPKHKEKMKIGVSPKOLGGRHESGYDNDPGLNLYVFCVDVKE 300
Db 241 IGMDDYVALLLEKPKHKEKMKIGVSPKOLGGRHESGYDNDPGLNLYVFCVDVKE 300
QY 301 TYDLLVQCCDAQPCASGYGVYVVMKRRQKQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLVQCCDAQPCASGYGVYVVMKRRQKQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPKVAQICYWKIGNYLDREG 383
Db 361 ITPKVAQICYWKIGNYLDREG 383

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RESULT 12

US-09-906-838-261
; Sequence 261, Application US/09906838
; Publication No. US20030027143A1
; GENERAL INFORMATION:

; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.

```

; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/906,838
; CURRENT FILING DATE: 2001-07-16
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-906-838-261

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Query Match 96.8%; Score 2044; DB 10; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPGCHKTGTPPTYKEAKOYLSYETLYANGSRTEXOVGIYILSSSGDGAXXRDSGS 120
Db 61 VSSSCGPGCHKTGTPPTYKEAKOYLSYETLYANGSRTEXOVGIYILSSSGDGAXXRDSGS 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
QY 181 KTVYGTQKLRVGLFKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTHTVPGKWIKNAND 240

Db 181 KTYVGTQKLRVGLFKPKFKDGGRGANDSTSAMPEQMKFQWIRVXRTHVPGKWKGNAND 240
Qy 241 IGWDYDVALLELKKPKRKFVKMGVSPAKQLPGGRTHFSGYDNDPRGNLVYRFCVDKDE 300
Db 241 IGWDYDVALLELKKPKRKFVKMGVSPAKQLPGGRTHFSGYDNDPRGNLVYRFCVDKDE 300
Qy 301 TYDLLYQQCDAQPGASGYVYVMWKRQKWKERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAQPGASGYVYVMWKRQKWKERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Qy 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 13

US-09-907-613-261
; Sequence 261, Application US/09907613
; Publication No. US20030027145A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, David
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavlin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,613
; CURRENT FILING DATE: 2001-07-17
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR FILING DATE: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564

; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-907-613-261
Query Match 96.8%; Score 2044; DB 10; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;
Qy 1 MAGIFGLLELFLFLLCAVQNSPYSAPWKPTWPAYRLPVVLPOSTLNLAKPDEGAEKLE 60
Db 1 MAGIFGLLELFLFLLCAVQNSPYSAPWKPTWPAYRLPVVLPOSTLNLAKPDEGAEKLE 60
Qy 61 VSSSCGPOCHKGTPLPTYKEAKQYLSYETLYANGSRTEXQVGIYILSSSGDGAXXRDSGS 120
Db 61 VSSSCGPOCHKGTPLPTYEEAKQYLSYETLYANGSRTEXQVGIYILSSSGDGAXXRDSGS 120
Qy 121 SGKRRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGCTCTLVAXKHVLTAAHCHD 180
Db 121 SGKRRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGCTCTLVAXKHVLTAAHCHD 180
Qy 181 KTYVGTQKLRVGLFKPKFKDGGRGANDSTSAMPEQMKFQWIRVXRTHVPGKWKGNAND 240
Db 181 KTYVGTQKLRVGLFKPKFKDGGRGANDSTSAMPEQMKFQWIRVXRTHVPGKWKGNAND 240
Qy 241 IGWDYDVALLELKKPKRKFVKMGVSPAKQLPGGRTHFSGYDNDPRGNLVYRFCVDKDE 300
Db 241 IGWDYDVALLELKKPKRKFVKMGVSPAKQLPGGRTHFSGYDNDPRGNLVYRFCVDKDE 300
Qy 301 TYDLLYQQCDAQPGASGYVYVMWKRQKWKERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAQPGASGYVYVMWKRQKWKERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Qy 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 14

US-09-907-942-261
; Sequence 261, Application US/09907942
; Publication No. US20030027146A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, David
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavlin, Ivar J.

APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/907,942
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-907-942-261

Query Match 96.8%; Score 2044; DB 10; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFFLLCAVGVSPYSAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFFLLCAVGVSPYSAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

Qy 61 VSSSCGQCHKGTPLPYKAKQVLSYETLYANGSRTEXOVGIYILSSGSGDGAAXRDSGS 120
Db 61 VSSSCGQCHKGTPLPYKAKQVLSYETLYANGSRTEXOVGIYILSSGSGDGAAXRDSGS 120

Qy 121 SGKRRKQIYGVDSRESIFGKQFLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180
Db 121 SGKRRKQIYGVDSRESIFGKQFLNYPSTSVKLSGTCTGLVAEXHVLTAACHIDG 180

Qy 181 KTVYKGTQKLRVGLKPKFKDGGKRGANDSAMPQMKFQWIRKRVTHVPKGIKGNAND 240
Db 181 KTVYKGTQKLRVGLKPKFKDGGKRGANDSAMPQMKFQWIRKRVTHVPKGIKGNAND 240

Qy 241 IGMDYDVALLELKKPHKRFKMGKIVGSPPAQKLPGRTHFSGYDNDRGNLVYRFDVKDE 300
Db 241 IGMDYDVALLELKKPHKRFKMGKIVGSPPAQKLPGRTHFSGYDNDRGNLVYRFDVKDE 300

Qy 301 TYDLLYQQCCDAQPGASGYGVYVRMWRQKQKWERKLIIGFSGHQWVDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCCDAQPGASGYGVYVRMWRQKQKWERKLIIGFSGHQWVDMNGSPQDFNVAVR 360

Qy 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 15
US-09-904-859-261
Sequence 261, Application US/09904859
Publication No. US20030036060A1
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Deenoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/904,859
CURRENT FILING DATE: 2001-07-12
PRIOR APPLICATION NUMBER: 09/665,350
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30

; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-904-859-261

Query Match 96.8%; Score 2044; DB 10; Length 383;
Best Local Similarity 98.4%; Pred. No. 4e-193;
Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

Qy 1 MAGTGGLLFLFLCAVGVSPYSAPWKPTWPAAYRLPVVLPOSTLNIAKPDGAEAKLE 60
Db ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
1 MAGTGGLLFLFLCAVGVSPYSAPWKPTWPAAYRLPVVLPOSTLNIAKPDGAEAKLE 60

Qy 61 VSSSCGPGCHKGTPPTYKEAKQVLSYETLYANGSRTEXQVGIYIILSSGDGAXXRDSDS 120
Db ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
61 VSSSCGPGCHKGTPPTYEAKQVLSYETLYANGSRTEXQVGIYIILSSGDGAXXRDSDS 120

Qy 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSTGCTGLVAEXHVLTAACHIDG 180
Db ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSTGCTGLVAEXHVLTAACHIDG 180

Qy 181 KTYVKGTKLVRGFLKPKFYDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
Db ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
181 KTYVKGTKLVRGFLKPKFYDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240

Qy 241 IGMDYDVALLELKKPHKPKFMKIGVSPPAKQLPGGRIFHSGYDNDPGLVYRFCVDKDE 300
Db ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
241 IGMDYDVALLELKKPHKPKFMKIGVSPPAKQLPGGRIFHSGYDNDPGLVYRFCVDKDE 300

Qy 301 TYDLLYQQCDAQPGASGYVYVRMWKRQQQKWERKIIGIPSGHQWYDMNGSPQDFNVAVR 360
Db ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
301 TYDLLYQQCDAQPGASGYVYVRMWKRQQQKWERKIIGIPSGHQWYDMNGSPQDFNVAVR 360

Qy 361 ITPLKYAQICYWIKGNVLDREG 383
Db ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
361 ITPLKYAQICYWIKGNVLDREG 383

Search completed: July 1, 2005, 21:32:16
Job time : 93.3582 secs

GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: July 1, 2005, 20:54:23 ; Search time 20.826 Seconds
(without alignments)
1811.048 Million cell updates/sec

Title: US-09-658-677-2

Perfect score: 2112

Sequence: 1 MAGIPGLFLRLLCAVQ.....IKGNLYDCREGDTVFLPGSN 392

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

- 1: PIR1:*
- 2: PIR2:*
- 3: PIR3:*
- 4: PIR4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	185	8.8	316	2 A45134	endopeptidase (EC
2	130.5	6.2	313	2 A35122	metalloproteinase
3	125	5.9	482	1 EXRT	coagulation factor
4	124	5.9	218	2 E97915	choline binding pr
5	118	5.6	269	2 B26823	pancreatic elastas
6	114	5.4	288	4 S70439	pancreatic elastas
7	114	5.4	287	4 A56615	probable pancreati
8	114	5.4	488	1 EXHU	coagulation factor
9	114	5.4	492	1 EXBO	coagulation factor
10	113	5.4	269	2 A26823	pancreatic elastas
11	113	5.4	271	2 A25528	pancreatic elastas
12	112.5	5.3	289	1 TRSGM	trypsin (EC 3.4.21
13	112	5.3	266	1 ELPG	pancreatic elastas
14	112	5.3	266	1 ELRT1	pancreatic elastas
15	112	5.3	522	2 T29767	hypothetical prote
16	111.5	5.3	238	1 TRW5Y	trypsin-like prote
17	111.5	5.3	686	1 A59271	Ra-reactive factor
18	111	5.3	246	1 DBHU	complement factor
19	110	5.2	269	2 C26823	pancreatic elastas
20	109	5.2	761	2 JC5759	brain-specific ser
21	107.5	5.1	285	2 C95045	choline binding pr
22	107.5	5.1	405	2 T35117	probable secreted
23	106	5.0	236	2 A28566	T-cell suppressor
24	106	5.0	271	1 ELRT2	pancreatic elastas
25	104.5	4.9	273	2 E85765	hypothetical prote
26	104.5	4.9	273	2 H64915	Putative protease
27	104.5	4.9	1582	2 T15308	hypothetical prote
28	102.5	4.9	2055	2 T31617	hypothetical prote
29	102	4.8	786	1 A47547	serine proteinase

30	101.5	4.8	278	2 AH0282	probable papetidas
31	101.5	4.8	583	2 A29154	complement factor
32	101.5	4.8	747	2 I51579	complement factor
33	101	4.8	1047	2 A55617	masquerade precurs
34	99	4.7	274	2 S40004	trypsin-related pr
35	97.5	4.6	259	2 S68424	allergen Der f III
36	97.5	4.6	272	2 JC4170	trypsin-like prote
37	96	4.5	407	1 KFB07	coagulation factor
38	94.5	4.5	409	2 T35118	probable secreted
39	94.5	4.5	416	1 KFB0	coagulation factor
40	94.5	4.5	1238	2 T34929	hypothetical prote
41	94.5	4.5	2145	2 JC4747	adenylate cyclase
42	94	4.5	266	2 JC4850	trypsin-like prote
43	94	4.5	624	2 T02289	probable polygalac
44	94	4.5	855	2 JC7731	membrane-bound arg
45	93.5	4.4	452	1 A30351	coagulation factor

ALIGNMENTS

RESULT 1

A45134

endopeptidase (EC 3.4.-.-), glutamate-specific - Bacillus licheniformis

C:Species: Bacillus licheniformis

C>Date: 10-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 16-Aug-2004

C/Accession: A45134; S23078

R;Kakudo, S.; Kikuchi, N.; Kitadokoro, K.; Fujiwara, T.; Nakamura, E.; Okamoto, H.; Shin,

J. Biol. Chem. 267, 23782-23788, 1992

A>Title: Purification, characterization, cloning, and expression of a glutamic acid-speci

A/Reference number: A45134; MUID:93054737; PMID:1429718

A/Accession: A45134

A>Status: preliminary

A/Molecule type: DNA

A/Residues: 1-316 <KX>

A/Cross-references: UNIPROT:P80057; GB:D10060; NID:G216263; PIDN:BAA00949.1; PID:D100141;

A/Experimental source: ATCC 14580

A/Note: sequence extracted from NCBI backbone (NCBIN:118784, NCBIP:118785)

R;Svendsen, I.; Breddam, K.

Eur. J. Biochem. 204, 165-171, 1992

A>Title: Isolation and amino acid sequence of a glutamic acid specific endopeptidase from

A/Reference number: S23078; MUID:92155199; PMID:1346764

A/Accession: S23078

A>Status: preliminary

A/Molecule type: protein

A/Residues: 95-316 <SV5>

C:Superfamily: Glutamyl endopeptidase, V8 type

C/Keywords: hydrolase

Query Match 8.8%; Score 185; DB 2; Length 316;

Best Local Similarity 23.2%; Pred. No. 4.2e-08;

Matches 77; Conservative 46; Mismatches 139; Indels 70; Gaps 16;

QY 63 SSCGQCHKGTPL---PTVKEAKQVLYETLYANGSRTEKXQVGYLSSSGDGAAXRDG 119

Db 28 AQAAPSPH--TPVSSDPSYK-AETSVYDP-----NIKSDQGLYSKAFITG---KVN 76

QY 120 SSGKRRK-----ROIYGYDSRFSIFGKDFLLNYPFSTSVKLST---GCTGLVAE 167

Db 77 TKEAEKSPAKAPYSIKSVIGSDRTRVN---TTAPYRAIVHSSIGSGCTGMIGP 133

QY 168 XHVLTAACIHGKGT-YVKGTKLRVGLKPKFKDGGGRANDSTAMPQMKFQWTRVKR 226

Db 134 KTVATAGHCIDYTSSTSGSPAGTATVSPG-----RNGTS-----YPYGSVKSTR 175

QY 227 THVPKGMKGNANDIGMDYDALLBELKPKHKFKMKIYGVSPPAKOLPGCRIHFSGVDNDR 286

Db 176 YFIPSGWRSGNTN-----YDYGAIELSEPIGNTVGYFGYSYTTSSLVGTVTISGYPGDK 230

QY 287 PGNLVVRFCDD--VKDETVDLLYQQCDAQPGASGYGVYVMMKRRQOKWKERKII---GIF 340

Db 231 TAGTQWQHSGPTAISEYTKLOYAM-DTYGGQSGSPVFEQSSSRSTNCSGCSLAVHTNGVY 289

QY 341 SGHWDVMSQDFNVAVRITPLKVAQICW 372
Db 290 GG-----SSYNRGTRITKEVFDNLTNW 311

RESULT 2
A35122
metalloprotease (EC 3.4.21.6) mpr precursor, extracellular - Bacillus subtilis
C:Species: Bacillus subtilis
C>Date: 27-Jul-1990 #sequence revision 27-Jul-1990 #text_change 16-Aug-2004
R:Accession: A35122; I40010; A69660
C:Sloma, A.; Rudolph, C.F.; Rufo Jr., G.A.; Sullivan, B.J.; Theriault, K.A.; Ally, D.; J. Bacteriol. 172, 1024-1029, 1990
A:Title: Gene encoding a novel extracellular metalloprotease in Bacillus subtilis.
A:Reference number: A35122; MUID:90130256; PMID:2105291
A:Accession: A35122
A>Status: preliminary
A:Molecule type: DNA
A:Residues: 1-313 <SLO>
A:Cross-references: UNIPROT:P39790; GB:L10505; NID:g143209; PIDN:AAA22604.1; PID:g143210
R:Smith, H.; de Jong, A.; Bron, S.; Venema, G.
Gene 70, 351-361, 1988
A:Title: Characterization of signal-sequence-coding regions selected from the Bacillus subtilis
A:Reference number: I39394; MUID:89108019; PMID:3145906
A:Accession: I40010
A>Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: DNA
A:Residues: 1-60, 65, 'L', 67, 'S', 69, 'AQA' <RES>
A:Cross-references: GB:M22916; NID:g143701; PIDN:AAA22832.1; PID:g143702
R:Kunst, F.; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Berte
C.; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, J.; Carter, N.M.; Ch
A.; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
Nature 390, 249-256, 1997
A:Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Gall
iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Hleapfel, S.; Hosono, S.; Hullo, M.F.
Koetter, P.; Konigstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois,
Y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetelle
Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon,
A:Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Seron
akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchiyama
T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, K
A:Authors: Yoshikawa, H.F.; Zumschein, E.; Yoshikawa, H.; Danchin, A.
A:Title: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis.
A:Reference number: A69580; MUID:98044033; PMID:93384377
A:Accession: A69660
A>Status: preliminary; nucleic acid sequence not shown; translation not shown
A:Molecule type: DNA
A:Residues: 1-313 <KUN>
A:Cross-references: GB:299105; GB:AL009126; NID:g2632457; PIDN:CAB12018.1; PID:el182176
A:Experimental source: strain 168
C:Genetics:
A:Gene: mpr
C:Superfamily: Glutamyl endopeptidase, V8 type
C:Keywords: hydrolase

Query Match 6.2%; Score 130.5; DB 2; Length 313;
Best Local Similarity 22.6%; Pred. No. 0.0018;
Matches 87; Conservative 38; Mismatches 139; Indels 121; Gaps 19;

QY 28 WKPTWPAFLPVVLPSTLNLAKPDFGAERAKLEVS-----SSCGPOCHKGTPLPTVYKEAK 82
Db 7 FRKQWFAYLTVLCLALA---AAVSFGVPAKAENPQTSVSNTKGEA-DATKNQTSKADQ 61

QY 83 QYLSYE-----TLVANGSRTEQXGVYILSSGDGAXXRDSGSGSKRRKQIYGYDS 135
Db 62 VSAPIYSGTGKTSKSLY--GGQTELEKNIOQLPS-----SIIGTDE 100

QY 136 RFSIFGKDFLLNYPFSTSVKLSL-----CCTGTLVAEXHVLTAACHIH----- 178
Db 101 RTRI-----SSSTSPFYPATVQLSIKYKPNSTSYCTGFLNPNVTVTAGHCVVYSDHGWAS 157

QY 179 -----DGKTVYKGTQKLRVGLKPKFKDGGGRANDSTSAMPEQMKFWIRVKRTHVP 230

Db 158 TITAPGRNGSSYPYGT-----SGTMFYSVK--GWTESKDTNVD 195
QY 231 KGWIKN---ANDIGWDYDYLLELKKPKRKFMKIGVSPPAKQLPGGRHFSGVDNDRP 287
Db 196 YCAIKLNGSPGNTVGW--GYRTTNSSP-----VGLSSSVTGPCKDTFGTMSDTPK 247
QY 288 GNLVRFCDKDETDYLLYQQCDQAGAGGVYVMKQKQKWERKIIGIFSGHQWVD 347
Db 248 IR-----SAETVKLTLY--TTDTYGCQSGSPVY-----RNYSDTGQTAIAIHT----- 287
QY 348 MNGSQDFNVAVRITPLKVAQICW 372
Db 288 -NGG-SSYNLGRTRVNDVFNFIQYW 310

RESULT 3
EXPT
coagulation factor Xa (EC 3.4.21.6) precursor - rat
C:Species: Rattus norvegicus (Norway rat)
C>Date: 31-Jan-1995 #sequence revision 07-Feb-1997 #text_change 09-Jul-2004
C:Accession: S49075; JC4670; PS0191; PS0190; I62745
R:Stanton, C.; Ross, P.; Hutson, S.; Wallin, R.
Thromb. Res. 80, 63-73, 1995
A:Title: Evidence for competition between vitamin K-dependent clotting factors for intrac
A:Reference number: A58498; MUID:96093366; PMID:8578539
A:Accession: S49075
A:Molecule type: mRNA
A:Residues: 1-482 <STA1>
A:Cross-references: UNIPROT:Q63207; EMBL:X79807; NID:g506600; PIDN:CAA56202.1; PID:g50660
A:Note: submitted to the EMBL Data Library, June 1994
A:Note: neither the complete nucleic acid sequence nor the complete translation are show
R:Stanton, C.; Ross, R.P.; Hutson, S.; Wallin, R.
Gene 169, 269-273, 1996
A:Title: Processing and expression of rat and human clotting factor-X-encoding cDNAs.
A:Reference number: JC4670; MUID:96194815; PMID:8647460
A:Accession: JC4670
A:Molecule type: mRNA
A:Residues: 1-482 <STA2>
A:Cross-references: EMBL:X79807; NID:g506600; PIDN:CAA56202.1; PID:g5066001
A:Experimental source: Cos-1 cell
R:Enjyoji, K.; Miyazaki, K.; Kato, H.
J. Biochem. 109, 890-898, 1991
A:Title: Characterization of rat factors X and Xa: demonstration of factor Xa in rat plas
A:Reference number: PS0190; MUID:92041742; PMID:1718949
A:Accession: PS0191
A:Molecule type: protein
A:Residues: 41-58, 'X', 60-65 <ENJ1>
A:Accession: PS0190
A:Molecule type: protein
A:Residues: 183-186, 'X', 188-207 <ENJ2>
R:Murakawa, M.; Okamura, T.; Kamura, T.; Kuroiwa, M.; Harada, M.; Niho, Y.
Eur. J. Haematol. 52, 162-168, 1994
A:Title: Analysis of the partial nucleotide sequences and deduced primary structures of t
A:Reference number: I46196; MUID:94222160; PMID:8168596
A:Accession: I62745
A>Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: DNA
A:Residues: 295-383, 'G', 385-455 <MUR>
A:Cross-references: GB:D21215; NID:g415309; PIDN:BA04756.1; PID:g455396
C:Function:
A:Description: catalyzes the proteolytic activation of prothrombin to thrombin in the pre
A:Pathway: blood coagulation
C:Superfamily: coagulation factor X; EGF homology; Gla domain homology; tryptsin homology
C:Keywords: beta-hydroxyaspartic acid; blood coagulation; calcium binding; carboxylglutami
F:1-23/Domain: signal sequence #status predicted <SIG>
F:25-84/Domain: propeptide #status predicted <PRO>
F:41-179/Product: coagulation factor X light chain #status predicted <LCH>
F:129-164/Domain: EGF homology <EG1>
F:183-482/Product: coagulation factor X heavy chain #status predicted <HCH>
F:183-231/Domain: activation peptide #status predicted <APT>


```

QY 358 AVRI 361
Db 202 AVKL 205

RESULT 5
B26823
pancreatic elastase II (EC 3.4.21.71) A precursor - human
C:Species: Homo sapiens (man)
C:Date: 16-Aug-1988 #sequence revision 16-Aug-1988 #text_change 09-Jul-2004
C:Accession: B26823; A27432; A14131; S34491
R:Kawashima, I.; Tani, T.; Shimoda, K.; Takiguchi, Y.
DNA 6, 163-172, 1987
A:Title: Characterization of pancreatic elastase II cDNAs: two elastase II mRNAs are expressed
A:Reference number: A9058; MUID:872117962; PMID:3646943
A:Accession: B26823
A:Molecule type: mRNA
A:Residues: 1-269 <KAW>
A:Cross-references: UNIPROT:P08217; GB:M16652; NID:g182057; PIDN:AAAS2380.1; PID:g182058
R:Fletcher, T.S.; Shen, W.F.; Largman, C.
Biochemistry 26, 7256-7261, 1987
A:Title: Primary structure of human pancreatic elastase 2 determined by sequence analysis
A:Reference number: A27432; MUID:88107669; PMID:3427074
A:Accession: A27432
A:Molecule type: mRNA
A:Residues: 1-269 <GB>
A:Cross-references: NID:g182022; PIDN:AAAS2374.1; PID:g182023
R:Shirasu, Y.; Yoshida, H.; Matsuki, S.; Takemura, K.; Ikeda, N.; Shimada, Y.; Ozawa, T.
J. Biochem. 102, 1555-1563, 1987
A:Title: Molecular cloning and expression in Escherichia coli of a cDNA encoding human pancreatic elastase
A:Reference number: A14131; MUID:88198076; PMID:2834346
A:Accession: A14131
A:Molecule type: mRNA
A:Residues: 1-201, 'V', 203-269 <SHI>
A:Cross-references: GB:D00336; NID:g219619; PIDN:BA00165.1; PID:g219620
A:Note: the authors translated the codon GTG for residue 202 as Cys
R:Moulard, M.; Michon, T.; Kerfelec, B.; Chapus, C.
FEBS Lett. 261, 179-183, 1990
A:Title: Further studies on the human pancreatic binary complexes involving procarboxypeptidase
A:Reference number: S08253; MUID:90169111; PMID:2307232
A:Accession: S34491
A:Molecule type: protein
A:Residues: 'X', 18-50 <MOU>
C:Genetics:
A:Gene: GDB:ELAI
A:Cross-references: GDB:119866; OMIM:130120
A:Map position: 12pter-12qter
C:Superfamily: trypsin; trypsin homology
C:Keywords: hydrolase; pancreas; serine proteinase
F:16/Domain: signal sequence #status predicted <SIG>
F:17-28/Domain: propeptide #status predicted <PRO>
F:29-269/Product: pancreatic elastase IIA #status predicted <MAT>
F:29-262/Domain: trypsin homology <TRY>
F:73,121,216/Active site: His, Asp, Ser #status predicted

Query Match 5.68; Score 118; DB 2; Length 269;
Best Local Similarity 24.1%; Pred. No. 0.038;
Matches 62; Conservative 24; Mismatches 67; Indels 104; Gaps 14;

QY 147 NYPFSTSVKLSTG-----CTGTLVAEKHVLTAACHIDGKTYVKGTKLRVGFLEKPKFK 200
Db 39 SWPQVSLQVSSNGKWHTCGSLIANSWLVLTAAHCISSTRY-----RVGL----- 85
QY 201 DGRG-----ANDSTSAMPEQKFWIRVKTHTVPKGTWTKGNANDIGMDYDVALLEKKP- 255
Db 86 --GRHNLVAESGSLA-----VSVSKTVVHKDW---NSNQISKGNDAIALLKLANPV 131
QY 256 HKKEFKMIGVSPAKQLPGGRIHFSGYDNDPGLVYRPFCDVKDEYDLLYQOCDAPGA 315
Db 132 SLTDKIQACLPAA-----GTI-----LPNN----- 152
QY 316 SGYGVTVRMKQOQKWERKIIGFSGHQWDMNGSPQDFNVAVRITPLKYA---QICW 372

```


Eur. J. Biochem. 218, 153-163, 1993
A:Title: Identification of O-linked oligosaccharide chains in the activation peptides of
A:Reference number: S39414; MUID:94062825; PMID:8243461
A:Accession: S39415
A:Molecule type: protein
A:Residues: 183-234 <INO>
A:Note: glycosylation sites
R:Jagadeeswaran, P.; Reddy, S.V.; Rao, K.J.; Hamsabhuhanam, K.; Lyman, G.
Gene 84, 517-519, 1989
A:Title: Cloning and characterization of the 5' end (exon 1) of the gene encoding human
A:Reference number: I54051; MUID:90128299; PMID:2612918
A:Accession: I54051
A:Status: translation not shown; translated from GB/ENBL/DBJ
A:Molecule type: DNA
A:Residues: 1-23 <RES>
A:Cross-references: GB:M33297; NID:g183860; PIDN:AAA52636.1; PID:g553330
R:Padmanabhan, K.; Padmanabhan, K.P.; Tulinsky, A.; Park, C.H.; Bode, W.; Huber, R.; Bla
J. Mol. Biol. 232, 947-966, 1993
A:Title: Structure of human des(1-45) factor Xa at 2.2 angstroms resolution.
A:Reference number: A49458; MUID:93360277; PMID:8355279
A:Contents: annotation; X-ray crystallography, 2.2 angstroms
C:Comment: The two chains held together by one disulfide bond are formed from a single-c
C:Comment: The activation peptide is cleaved by factor IXa (in the intrinsic pathway) or
C:Genetics:
A:Gene: GDB:F10
A:Cross-references: GDB:119890; OMIM:227600
A:Map position: 13q34-13q34
A:Introns: 24/1; 77/3; 86/1; 124/1; 150/3; 249/3; 289/1
A:Note: deficiency of this factor causes Stuart disease
C:Function:
A:Description: catalyzes the proteolytic activation of prothrombin to thrombin in the pr
A:Pathway: blood coagulation
C:Superfamily: coagulation factor X; EGF homology; Gla domain homology; trypsin homology
C:Keywords: beta-hydroxyaspartic acid; blood coagulation; calcium binding; carboxyglutam
F:1-23/Domain: signal sequence #status predicted <SIG>
F:24-40/Domain: propeptide #status predicted <PRO>
F:25-84/Domain: Gla domain homology <GLA>
F:41-179/Product: coagulation factor X light chain #status experimental <LCH>
F:129-161/Domain: EGF homology <EG1>
F:183-488/Product: coagulation factor X heavy chain #status experimental <HCH>
F:183-234/Domain: activation peptide #status experimental <APT>
F:235-488/Product: coagulation factor Xa heavy chain #status experimental <ACT>
F:235-462/Domain: trypsin homology <TRY>
F:46,47,54,56,59,60,65,66,69,72,79/Modified site: gamma-carboxyglutamic acid (Glu) #stat
F:57-62/Disulfide bonds: #status predicted
F:90-101,95-110,112-121,129-140,136-149,151-164,172-342,241-246,261-277,390-404,415-443/
F:103/Modified site: erythro-beta-hydroxyaspartic acid (Asp) #status experimental
F:199,211/Binding site: carboxylate (Thr) (covalent) #status experimental
F:221,231/Binding site: carboxylate (Asn) (covalent) #status experimental
F:234-235/Cleavage site: Arg-Ile (coagulation factor IXa, coagulation factor VIIa) #stat
F:276,322,419/Active site: His, Asp, Ser #status experimental

Query Match 5.4%; Score 114; DB 1; Length 488;
Best Local Similarity 20.9%; Pred. No. 0.08;
Matches 57; Conservative 37; Mismatches 67; Indels 112; Gaps 13;
QY 160 CTGLVAEXHLVTAACHIDHDKTYVKGTKLVGFLKPKFGDGGANDSTAMPQMKF 219
DB 261 CGTILSEFILTANCLVQAKF-----KVRVGRDNTQEEGEAVHE-VEVVIKHNFP 314
QY 220 QWIRVTRTHVPKGIWKNANDIGMDYDYLLELLKPKHKKFMKIGVSPPA----- 269
DB 315 -----TKETV-----DFDIIVLRKLTPTIT---FRMNVAPACLPERDWAEST 352
QY 270 --KQLPG-----CRIHFGVDNDPGLNLYRFDVKD-----ETVDLLVQ 307
DB 353 LMTQTKTGVSGFGRTEKGRQSLKMLKLEVPYVYDRNSCKLSSSFITQNMFCAGYDT--K 410
QY 308 QCAQAPCASG-----YGVYVRMKRQOQKWERKIIGI 339
DB 411 QEDACQDGGPHTVRFKDTYFTGIVSGWEGCARKGKGIYTK-----V 455

QY 340 FSGHWDVDMN-----GSPDENVAVRI---TPLK 365
DB 456 TAPLKWIDRSMKTRGLPKAKSHAPEVITSSPLK 488

RESULT 9

EXPO

N:Alternates names: Stuart factor
N:coagulation factor Xa (EC 3.4.21.6) precursor - bovine
C:Species: Bos primigenius taurus (cattle)
C:Date: 24-Apr-1984 #sequence revision 17-Mar-1987 #text change 09-Jul-2004
C:Accession: A22867; A14997; A12030; A34412; S39414; A00925
R:Pung, M.R.; Campbell, R.M.; MacGillivray, T.A.
Nucleic Acids Res. 12, 4481-4492, 1984
A:Title: Blood coagulation factor X mRNA encodes a single polypeptide chain containing a
A:Reference number: A22867; MUID:84247315; PMID:6330671
A:Accession: A22867
A:Molecule type: mRNA
A:Residues: 1-487 <FUN>
A:Cross-references: UNIPROT:P00743; GB:X00673; NID:g192; PIDN:CAA25286.1; PID:g193
R:Enfield, D.L.; Ericsson, L.H.; Fujikawa, K.; Walsh, K.A.; Neurath, H.; Titani, K.
Biochemistry 19, 659-667, 1980
A:Title: Amino acid sequence of the light chain of bovine factor X-1 (Stuart factor).
A:Reference number: A14997; MUID:80130563; PMID:6766735
A:Accession: A14997
A:Molecule type: protein
A:Residues: 41-102, 'N', 104-180 <ENF>
R:McMullen, B.A.; Fujikawa, K.; Kisiel, W.
Biochem. Biophys. Res. Commun. 115, 8-14, 1983
A:Title: The occurrence of beta-hydroxyaspartic acid in the vitamin K-dependent blood co
A:Reference number: A20274; MUID:83308813; PMID:6688526
A:Contents: annotation; revision to residue 103
R:Titani, K.; Fujikawa, K.; Enfield, D.L.; Ericsson, L.H.; Walsh, K.A.; Neurath, H.
Proc. Natl. Acad. Sci. U.S.A. 72, 3082-3086, 1975
A:Title: Bovine factor X-1 (Stuart factor): amino-acid sequence of heavy chain.
A:Reference number: A12030; MUID:76053069; PMID:1059093
A:Accession: A12030
A:Molecule type: protein
A:Residues: 183-292,294-295, 'GDE', 299-334,336-348, 'AB', 351-354,356-441, 'QKFG', 446-492 <TJ
A:Note: carboxylate binding sites and disulfide bonds were determined
R:Persson, E.; Selander, M.; Linse, S.; Drakenberg, T.; Oehlin, A.K.; Stenflo, J.
J. Biol. Chem. 264, 16897-16904, 1989
A:Title: Calcium binding to the isolated beta-hydroxyaspartic acid-containing epidermal s
A:Reference number: A34412; MUID:89380326; PMID:2789221
A:Accession: A34412
A:Molecule type: protein
A:Residues: 85-126 <PER>
A:Note: beta-hydroxyaspartic acid site
R:Inoue, K.; Morita, T.
Eur. J. Biochem. 218, 153-163, 1993
A:Title: Identification of O-linked oligosaccharide chains in the activation peptides of
A:Reference number: S39414; MUID:94062825; PMID:8243461
A:Accession: S39414
A:Molecule type: protein
A:Residues: 183-196;199-209;216-233 <INO>
A:Note: carboxylate binding sites
R:Titani, K.; Hermanson, M.A.; Fujikawa, K.; Ericsson, L.H.; Walsh, K.A.; Neurath, H.; Dr
Biochemistry 11, 4899-4903, 1972
A:Title: Bovine factor X-1a (activated Stuart factor). Evidence of homology with mammalia
A:Reference number: A12453; MUID:73053314; PMID:4264286
A:Contents: annotation; active site
R:Fujikawa, K.; Titani, K.; Davie, E.W.
Proc. Natl. Acad. Sci. U.S.A. 72, 3359-3363, 1975
A:Title: Activation of bovine factor X (Stuart factor): conversion of factor Xaalpha to i
A:Reference number: A13504; MUID:76053121; PMID:1059122
A:Contents: annotation; activation
R:Sugo, T.; Bjork, I.; Holmgren, A.; Stenflo, J.
J. Biol. Chem. 259, 5705-5710, 1984
A:Title: Calcium-binding properties of bovine factor X lacking the gamma-carboxyglutamic
A:Reference number: A38024; MUID:84185716; PMID:6546930
A:Contents: annotation; calcium binding
R:Morita, T.; Jackson, C.M.

```
A;Accession: A26823
A;Molecule type: mRNA
A;Residues: 1-269 <RAW>
A;/Cross-references: UNIPROT:P08419; GB:M16651; NID:g164441; PIDN:AAA31027.1; PID:g164442
C:/Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase
F;1-16/Domain: signal sequence #status predicted <SIG>
F;17-28/Domain: propeptide #status predicted <PRO>
F;29-269/Product: elastase II #status predicted <MAT>
F;29-262/Domain: trypsin homology <TRY>
F;73,121,216/Active site: His, Asp, Ser #status predicted
```

Query Match 5.4%; Score 113; DB 2; Length 269;
Best Local Similarity 27.1%; Pred.No. 0.048;
Matches 59; Conservative 28; Mismatches 65; Indels 66; Gaps 14;

QY 128 ROIYGVDSRFSFGKDFLLNYPFSTSVKL-STG-----CTGTIVAEXHVLTAHCIIHDGK 181
 :|::||:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:
Db :|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:

QY 182 TY--VKGTKLVRGVFLPKPFGDGRGANDSTSAMPEQMFKFWIRVKRTHPVPKWIKGNAN 239
 :|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:
Db :|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:-NSN 114
 80 TYRVVLGRHSLSL-----STNEPGSLA-----VKYSKLVVHQDW----

QY 240 DTGMVDYYALLELKXP-HKRKPWKIGVSPPAKO-LPG-----GRHFSGYND--R 286
 :|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:
Db :|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:

QY 287 PGNLYVRFCDVDETYDLLYOCDAPGASGYGVYRM 324
 ||:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:
Db |:||:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:-YATC-SKPGWGWGSTVTXNM 200
 175 QGOLL-----VVDD-----

RESULT 11

A25528
pancreatic elastase II (EC 3.4.21.71) precursor - mouse
C;Species: Mus musculus (house mouse)
C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 09-Jul-2004
C;Accession: A25528
R;Stevenson, B.J.; Hagenbuechle, O.; Wellauer, P.K.
Nucleic Acids Res. 14, 8307-8330, 1986
A;Title: Sequence organisation and transcriptional regulation of the mouse elastase II gene
A;Reference number: A93646; MUID:87066713; PMID:3641189
A;Accession: A25528
A;Molecule type: mRNA
A;Residues: 1-271 <STE>
A;/Cross-references: UNIPROT:P05208; GB:X04573; NID:g50825; PIDN:CAA28242.1; PID:g50826
C;/Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase
F;1-30/Domain: signal sequence #status predicted <SIG>
F;31-271/Product: pancreatic elastase II #status predicted <MAT>
F;31-264/Domain: trypsin homology <TRY>
F;75,123,218/Active site: His, Asp, Ser #status predicted

Query Match 5.4%; Score 113; DB 2; Length 271;
Best Local Similarity 29.0%; Pred.No. 0.048;
Matches 38; Conservative 22; Mismatches 37; Indels 34; Gaps 7;

QY 148 YPFSTSVK-LSTG-----CTGTIVAEXHVLTAHCIIHDGKY--VKGTOKLRVGFLKPKF 199
 :|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:
Db :|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:

QY 200 KDGGRGANDSTSAMPEQMFKFWIRVKRTHPKWIKGNADIGMDYDALLELKXP-HKR 258
 :|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:
Db :|:|:|:|:|:|:|:|:|:|:|:|:|:|:~::~:-VQVSKLVVHQRW---NSQNGVNGYDIALLI KLASPVTLIS 136
 93 --SNTPAGSAA-----

QY 259 KPWKIGVSPPA 269
 :|:|:|:|:|:~::~:
Db 137 KNITACLPAPA 147
 :|:|:|:~::~:

RESULT 12

TRSMG

trypsin (EC 3.4.21.4) precursor - Streptomyces griseus
C:Species: Streptomyces griseus
C>Date: 24-Apr-1984 #sequence_revision 12-May-1994 #text_change 09-Jul-2004
C:Accession: JQ1302; A00962
R:Kim, J.C.; Cha, S.H.; Jeong, S.T.; Oh, S.K.; Byun, S.M.
Biochem. Biophys. Res. Commun. 181, 707-713, 1991
A:Title: Molecular cloning and nucleotide sequence of Streptomyces griseus trypsin gene.
A:Reference number: JQ1302; MUID:92095977; PMID:1755852
A:Accession: JQ1302
A:Molecule type: DNA
A:Residues: 1-259 <XIM>
A:Cross-references: UNIPROT:P00775; GB:M64471
A:Experimental source: strain ATCC10137
R:Olafson, R.W.; Jucasek, L.; Carpenter, M.R.; Smillie, L.B.
Biochemistry 14, 1168-1177, 1975
A:Title: Amino acid sequence of Streptomyces griseus trypsin. Cyanogen bromide fragments
A:Reference number: A00962; MUID:75127940; PMID:804314
A:Accession: A00962
A:Molecule type: protein
A:Residues: 37-95,98-259 <OLA>
R:Read, R.J.; James, M.N.G.
J. Mol. Biol. 200, 523, 1988
A:Title: Refined crystal structure of Streptomyces griseus trypsin at 1.7 angstroms resolution
A:Reference number: A44574; MUID:88286735; PMID:3135412
A:Contents: annotation; X-ray crystallography, 1.7 angstroms
A:Note: residues 96-97 modeled as Gly-Ala
C:Genetics:
A:Gene: sprt
C:Superfamily: trypsin; trypsin homology
C:Keywords: hydrolase; serine proteinase
F:1-32/Domain: signal sequence #status predicted <SIG>
F:33-36/Domain: propeptide #status predicted <PRO>
F:37-258/Product: trypsin #status experimental <MAT>
F:37-252/Domain: trypsin homology <TRY>
F:58-74,177-192,204-233/Disulfide bonds: #status experimental
F:73,118,208/Active site: His, Asp, Ser #status experimental
Query Match 5.3%; Score 112.5; DB 1; Length 259;
Best Local Similarity 26.0%; Pred. No. 0.05;
Matches 39; Conservative 21; Mismatches 53; Indels 37; Gaps 5;
QY 148 YPFSTSVKLSGCTGLVAEXHVLTAACHTIDGKTVKQKLVGFLKPKFKDGGGAN 207
DB 48 PFP-MVRLSMGCGALYAQDIVLTAACHV-----SSGNN 81
QY 208 DS---TSAMPEQMKFQWIRKTHVPKGIKGNANDIGMDYDVALLELKKPKKPMKIG 264
DB 82 TSITATGGVVDLQSSAVKVRSTKVLQA-----PGYNGTGKDWALIKLAQPINQPTLKA 136
QY 265 VSPPAQLPGGRHFGSYNDREGNLYRFP 294
DB 137 TTTAYNQ---GTTVAGWGANGREGGQQRY 163
RESULT 13
ELPG
pancreatic elastase (EC 3.4.21.36) I precursor - pig
C:Species: Sus scrofa domestica (domestic pig)
C>Date: 24-Apr-1984 #sequence_revision 30-Sep-1990 #text_change 09-Jul-2004
C:Accession: JS0013; A26777; A10061; A00959
R:Shirau, Y.; Yoshida, H.; Mikayama, T.; Matsuki, S.; Tanaka, J.I.; Ikenaga, H.
J. Biochem. 99, 1707-1712, 1986
A:Title: Isolation and expression in Escherichia coli of a cDNA clone encoding porcine H
A:Reference number: A92005; MUID:86304235; PMID:3528137
A:Accession: JS0013
A:Molecule type: mRNA
A:Residues: 1-266 <SHI>
A:Cross-references: UNIPROT:P00772; GB:X04036; GB:D00070; GB:N00070; MUID:g1941; PIDN:CAA
R:Fan, T.; Kawashima, I.; Furukawa, H.; Ohmine, T.; Takiguchi, Y.
J. Biochem. 101, 591-599, 1987
A:Title: Characterization of a silent gene for human pancreatic elastase I: structure of
A:Reference number: A26777; MUID:87250343; PMID:3648024
A:Accession: A26777

A:Molecule type: mRNA
A:Residues: 1-125,'G',127-183,'L',185-266 <TAN>
A:Cross-references: GB:D00160; MUID:9217683; PIDN:BAA00118.1; PID:g217684
A:Note: the authors translated the codon GGG for residue 58 as Gln, GGC for residue 126 &
R:Shotton, D.M.; Hartley, B.S.
Biochem. J. 131, 643-675, 1973
A:Title: Evidence for the amino acid sequence of porcine pancreatic elastase.
A:Reference number: A90267; MUID:73229121; PMID:4578945
A:Accession: A10061
A:Molecule type: protein
A:Residues: 27-91,'N',93-203,'N',205-266 <SHO>
R:Shotton, D.M.; Hartley, B.S.
Nature 225, 811-816, 1970
A:Title: Three-dimensional structure of tosyl-elastase.
A:Reference number: A93160; MUID:70114044; PMID:5415110
A:Contents: annotation; X-ray crystallography, 3.5 angstroms; active site
C:Superfamily: trypsin; trypsin homology
C:Keywords: hydrolase; pancreas; serine proteinase; zymogen
F:1-16/Domain: signal sequence #status predicted <SIG>
F:17-26/Domain: activation peptide #status predicted <APT>
F:27-266/Product: elastase I #status experimental <MAT>
F:27-259/Domain: trypsin homology <TRY>
F:56-72,153-220,184-200,210-240/Disulfide bonds: #status experimental
F:71,119,214/Active site: His, Asp, Ser #status experimental
Query Match 5.3%; Score 112; DB 1; Length 266;
Best Local Similarity 27.3%; Pred. No. 0.057;
Matches 36; Conservative 21; Mismatches 41; Indels 34; Gaps 6;
QY 147 NYPFSTSVKLSG-----CTGTLVAEXHVLTAACHTIDGKTV--VKGTQKLVGFLKPK 198
DB 37 SWPSQISIQYRSGSSWAHTCGGTLIRQNWVMTAAHCVDELFRVVVGHNH----- 88
QY 199 FKDGGGANDSTSAMPEQMKFQWIRKTHVPKGIKGNANDIGMDYDVALLEL-KKPKH 257
DB 89 -----NQNDGTE-----QYVGQKIVVHPYV---NTDDVAAGYDIALRLAQSVTL 131
QY 258 RKFMKIGVSPPA 269
DB 132 NSYVQLGVLPR 143
RESULT 14
ELRTI
pancreatic elastase (EC 3.4.21.36) I precursor - rat
C:Species: Rattus norvegicus (Norway rat)
C>Date: 18-Aug-1982 #sequence_revision 18-Aug-1982 #text_change 09-Jul-2004
C:Accession: A00960; A20534
R:MacDonald, R.J.; Swift, G.H.; Quinto, C.; Swain, W.; Pictet, R.L.; Nikovits, W.; Rutten
Biochemistry 21, 1453-1463, 1982
A:Title: Primary structure of two distinct rat pancreatic preproelastases determined by
A:Reference number: A00960; MUID:82182967; PMID:6918221
A:Accession: A00960
A:Molecule type: mRNA
A:Residues: 1-266 <MAC>
A:Cross-references: UNIPROT:P00773; GB:V01234; MUID:956088; PIDN:CAA24544.1; PID:g56089
R:largman, C.
Biochemistry 22, 3763-3770, 1983
A:Title: Isolation and characterization of rat pancreatic elastase.
A:Reference number: A20534; MUID:84000385; PMID:6555050
A:Accession: A20534
A:Molecule type: protein
A:Residues: 17-37,'X',39-45 <LAR>
C:Superfamily: trypsin; trypsin homology
C:Keywords: hydrolase; pancreas; serine proteinase; zymogen
F:1-16/Domain: signal sequence #status predicted <SIG>
F:17-26/Domain: activation peptide #status predicted <APT>
F:27-266/Product: elastase I #status predicted <MPT>
F:27-259/Domain: trypsin homology <TRY>
F:71,119,214/Active site: His, Asp, Ser #status predicted
Query Match 5.3%; Score 112; DB 1; Length 266;
Best Local Similarity 26.7%; Pred. No. 0.057;

Search completed: July 1, 2005, 21:09:18
Job time : 24.826 secs

RESULT 15

RESOLUTION 13
T29767

hypothetical protein ZC581.6 - *Caenorhabditis elegans*

R; Waterston, B.; Gattung, S.; Le, T.T.

A:Description: The sequence of *C. elegans* cosmid ZC581

A;Reference number: Z20682

A;Status: preliminary; translated from GB/EMBL/DBJ

A;Residues: 1-522 <WAT>

A;CROSS-References: UNIPROT:U01771; EMBL:AF003134; PDB: AAB54144.1; GSPDB:G
A-Eynerimental source: strain Bristol N2: clone ZC581

C;Genetics:

A;Map position: 1

Z/VLE 1T/T7E 1T/TCC 1T/TTC 1P/ECE 1C/CCT 1S/OO :CHOTAMUHU

Query match	5.38;	Score 112;	DB 2;	Length 522;
Best local similarity	20.68;	Pred NC	0.13.	

Matches 95; Conservative 53; Mismatches 160; Indels 154; Gaps

QY 29 KPTWPAYRLPVLPQSTLNLA KPDFGAEAKLEVSS--CGPQCHKGTP--LPTYKEAK

Db 26 KPSNKASSAPSLRKSSSNPNKGTARSVSKSVPKSSAIPASPTVQKEVPPVEIEKKKEEK

83 -QVI-SVETI.YANGSPTEXOVGIYTI.SSSGNGAXYBPDSGSSGKSPBPOTYGYNDSPESTEG

[illegible]

Variable	Mean	Standard Deviation	Minimum	Maximum
Age	35.2	12.5	22	65
Gender	0.48	0.50	0	1
Marital Status	0.65	0.48	0	1
Education	12.8	2.1	9	16
Income	45000	15000	20000	80000
Health	0.72	0.45	0	1
Smoking	0.25	0.43	0	1
Alcohol	0.18	0.38	0	1
Exercise	0.35	0.48	0	1
Stress	0.60	0.49	0	1
Depression	0.22	0.42	0	1
Loneliness	0.30	0.46	0	1
Life Satisfaction	0.70	0.45	0	1
Overall Health	0.75	0.43	0	1

DB 146 KIDEMILLYWLRIFFSAKVYNGRDASQSEAPWSVFITYLYSKDEQSAITCTGTIVSPRHILL

Qy 173 AAHCI---HDGK-----TYVKGTKL--RVGFLKPKFKDGGRGANDST

Db 206 ATHCFAGQNRDGSWNLIETFDRSNCKDDDYVTNQEFLKRVFLSNK-----KGI

Qv 211 SAMPEOMKFOWIRVKRTHVPKGWIKGNANDIGMPY---DYALLEL-----

257 SPYRDEYITV INACSTWPT - - - - - ANBETKVT PPOCVTDRERATUTVI VEET TEESNTI/OSIVICU

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[illegible][illegible]

QY 293 ---RFCUVRDETYD-----LLYQQCDAQ-----PGASGYGVYVRMWKRQQQKWERKII

Db 362 YFFQARDITDKTTVACVVS LKILILNK TQASLNIS LKGD SGGAIADV-----KGKKTII

Qy 338 GIFS-----GHQWVDMNGSPQDFNVAVRITPLKYAQIC 370

Db 416 GVL SOTS COKRRGGNETMELYSSVGFKNOI----CKYTGIC 453

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: July 1, 2005, 20:53:37 ; Search time 92.7095 Seconds
(without alignments)
2165.204 Million cell updates/sec

Title: US-09-658-677-2

Perfect score: 2112

Sequence: 1 MAGIGLFLFLFLCAVGQ.....IKGNVLDRCBGDTVFLPGSN 392

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1612378 seqs, 512079187 residues

Total number of hits satisfying chosen parameters: 1612378

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Uniprot 03:*

1: uniprot_sprot:*

2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	2044	96.8	383	1 PS23 HUMAN	O95084 homo sapien
2	1881	89.1	383	2 O6AY61	O6AY61 rattus norv
3	1866.5	88.4	382	1 PS23 MOUSE	O95084 mus musculus
4	1860.5	88.1	382	2 O8BZS4	O8BZS4 mus musculus
5	1029	48.7	413	2 O9BQP6	O9BQP6 homo sapien
6	1028	48.7	413	2 O8N320	O8N320 homo sapien
7	1003.5	47.5	409	2 O8CQF9	O8CQF9 mus musculus
8	998.5	47.3	409	2 O8CQF5	O8CQF5 mus musculus
9	994.5	47.1	418	2 O6GML6	O6GML6 brachydanio
10	992.5	47.0	409	2 O8CQD6	O8CQD6 mus musculus
11	185	8.8	316	1 GSEP BACLI	P80057 bacillus li
12	185	8.8	316	2 O65NR6	O65NR6 bacillus li
13	182	8.6	450	2 O71YE5	O71YE5 listeria li
14	165.5	7.8	384	2 O98G17	O98G17 rhizobium l
15	156	7.4	271	2 O987W6	O987W6 rhizobium l
16	133.5	6.3	358	2 O931E7	O931E7 staphylococ
17	132.5	6.3	323	2 O73D54	O73D54 bacillus ce
18	132	6.2	267	2 O6DGW4	O6DGW4 brachydanio
19	130.5	6.2	313	1 MPR BACSU	P39790 bacillus su
20	130	6.2	439	2 O9GMD9	O9GMD9 ornithorhyn
21	129.5	6.1	321	2 O7NGB4	O7NGB4 gloeobacter
22	128.5	6.1	678	2 O9JJ58	O9JJ58 rattus norv
23	128.5	6.1	1322	2 O9NAT0	O9NAT0 anopheles g
24	128	6.1	266	2 O6AZC0	O6AZC0 brachydanio
25	125.5	5.9	339	2 O9QX91	O9QX91 rattus norv
26	125.5	5.9	366	2 O9QX85	O9QX85 rattus norv
27	125.5	5.9	541	2 O9QX90	O9QX90 rattus norv
28	125.5	5.9	623	2 O9JJ53	O9JJ53 rattus norv
29	125.5	5.9	643	2 O9QX84	O9QX84 rattus norv
30	125.5	5.9	1234	2 O7P1Q7	O7P1Q7 anopheles g
31	125.5	5.9	1322	2 O7PNR7	O7PNR7 anopheles g

32	125.5	5.9	1322	2 O9NJS5	O9NJS5 anopheles g
33	125	5.9	482	1 FA10 RAT	O63207 rattus norv
34	124.5	5.9	290	2 O81HL5	O81HL5 bacillus ce
35	124	5.9	218	2 O8DR41	O8DR41 streptococc
36	123.5	5.8	376	1 FA10 HOPST	P83370 hoptocephal
37	123	5.8	259	2 O9XY61	O9XY61 ctienocephal
38	123	5.8	266	2 O46644	O46644 macaca fasc
39	123	5.8	352	2 O6UWB4	O6UWB4 homo sapien
40	122.5	5.8	303	2 O9EXR9	O9EXR9 bacillus in
41	122.5	5.8	490	1 FA10 RABIT	CI9045 oryctolagus
42	121	5.7	745	2 O9OWF9	O9OWF9 triakis scy
43	120	5.7	269	2 O6ICV2	O6ICV2 homo sapien
44	120	5.7	269	2 O6ISU5	O6ISU5 homo sapien
45	120	5.7	685	2 O91WP0	O91WP0 mus musculu

ALIGNMENTS

RESULT 1
PS23_HUMAN STANDARD; PRT; 383 AA.
AC O95084;
DT 28-FEB-2003 (Rel. 41, Created)
DT 28-FEB-2003 (Rel. 41, Last sequence update)
DT 25-OCT-2004 (Rel. 45, Last annotation update)
DE Serine protease 23 precursor (EC 3.4.21.-) (Putative secreted protein
DE ZSIG13) (UNQ270/PRO307).
GN Names=PRSS23; Synonym=ZSIG13;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Umbilical vein;
RA Li X., Tedder T.F.;
RT "A novel serine protease from human umbilical vein endothelial
RT cells.";
RL Submitted (JUL-1997) to the EMBL/GenBank/DBJ databases.
RN [2]
RP SEQUENCE FROM N.A.
RA Sheppard P., Blumberg H., Jelinek L., Foster D., O'Hara P.;
RL Submitted (OCT-1999) to the EMBL/GenBank/DBJ databases.
RN [3]
RP SEQUENCE FROM N.A.
RA TISSUE=Uterus;
RA MEDLINE=21154917; PubMed=11230166; DOI=10.1101/gr.154701;
RA Wiemann S., Weil B., Wellenreuther R., Gassenhuber J., Glaesl S.,
RA Ansoerge W., Boecker H., Bloeker H., Bauersachs S., Blum H.,
RA Lauber J., Dueterthof A., Beyer A., Koehrer K., Strack N.,
RA Mewes H.-W., Ottenwaelder B., Obermaier B., Tampe J., Heubner D.,
RA Wambutt R., Korn B., Klein M., Poustka A.;
RT "Towards a catalog of human genes and proteins: sequencing and
RT analysis of 500 novel complete protein coding human cDNAs.";
RL Genome Res. 11:422-435(2001).
RN [4]
RP SEQUENCE FROM N.A.
RA MEDLINE=22887296; PubMed=12975309; DOI=10.1101/gr.1293003;
RA Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brueh J.,
RA Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,
RA Eaton D., Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S.,
RA Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,
RA Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,
RA Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,
RA Vandlen R., Watanabe C., Wiedand D., Woods K., Xie M.-H., Yaneura D.,
RA Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,
RA Godowski P., Gray A.;
RT "The secreted protein discovery initiative (SPDI), a large-scale
RT effort to identify novel human secreted and transmembrane proteins: a
RT bioinformatics assessment.";
RL Genome Res. 13:2265-2270(2003).
RN [5]


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RESULT 3
PS23_MOUSE
ID PS23_MOUSE STANDARD; PRT; 382 AA.
AC Q9D6X6; Q8VEG1;
DT 28-FEB-2003 (Rel. 41, Created)
DT 28-FEB-2003 (Rel. 41, Last sequence update)
DT 25-OCT-2004 (Rel. 45, Last annotation update)
DE DE Serine protease 23 precursor (EC 3.4.21.-).
GN Name=Prs23;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mus.
OX NCBI_TaxID=10090;
RN [1]
SEQUENCE FROM N.A.
RP STRAIN=C57BL/6J; TISSUE=Tongue;
RC MEDLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;
RA Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,
RA Nikaïdo I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,
RA Vagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori T.,
RA Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
RA Schraml L.M., Knapin A., Matsuda H., Batalov S., Beisel K.W.,
RA Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA Dalla E., Dragani T.A., Fletcher C.P., Forrest A., Frazer K.S.,
RA Gaasterland T., Gariboldi M., Giasi C., Godzik A., Gough J.,
RA Grimmond S., Gustincich S., Hirokawa N., Jackson I.J., Jarvis E.D.,
RA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
RA Konagaya A., Kurochkin I.V., Lee Y., Lennard B., Lyons P.A.,
RA Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,
RA Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Resolle G.,
RA Petrovsky N., Pillai R., Pontius J.U., Qi D., Ramachandran S.,

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CC DR ENBL; BC018517; AAH18517.1; -.
CC DR HSSP; PC00746; IDSU.
CC DR MEROPS; S01.309; -.
CC DR MGD; MGI:1923703; 2310046G15Rik.
CC DR InterPro; IPR009003; Pept Ser Cys.
CC DR InterPro; IPR011254; Peptidase_S1.
CC DR InterPro; IPR001314; Peptidase_S1A.
CC DR Pfam; PF00089; Trypsin; 1.
CC DR PRINTS; PR00722; CHYMOTRYPSIN.
CC DR SMART; SM00020; Tryp_SPC; 1.
CC DR PROSITE; PS02400; TRYPSIN_DOM; FALSE_NEG.
CC DR PROSITE; PS00134; TRYPSIN_HIS; 1.
CC DR PROSITE; PS00135; TRYPSIN_SER; FALSE_NEG.
CC DR KWL HydroLase; Serine protease; Signal.
CC DR SIGNAL 1 22 Potential.
CC FT CHAIN 23 382 Serine protease 23.
CC FT ACT_SITE 174 174 Charge relay system (By similarity).
CC FT ACT_SITE 239 239 Charge relay system (By similarity).
CC FT ACT_SITE 315 315 Charge relay system (By similarity).
CC FT DISULFID 159 175 N-linked.
CC FT CARBOHYD 192 192 N-linked (GLCNAC... ) (Potential).
CC FT

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FT CARBOHYD 206 206 N-linked (GlcNAc...) (Potential).
SQ CONFLICT 260 260 M -> I (in Ref. 1; BAB26541).
SQ SEQUENCE 382 AA; 43071 MW; 6F09A5C80A5B2306 CRC64;

Query Match 88.4%; Score 1866.5; DB 1; Length 382;
Best Local Similarity 89.8%; Pred. No. 5.1e-157;
Matches 343; Conservative 14; Mismatches 24; Indels 1; Gaps 1;

QY 1 MAGIPGLFLLFLLCAVGQVSPYAPWKPPTWPAVRLPVVLPQSTLNLAQDFGABAKLE 60
DB 1 MAGIPG-LFILLVLLCVFMQVSPYTPWKPPTWPAVRLPVVLPQSTLNLAQDFGABAKLE 59

QY 61 VSSSCGPGCHKGTPLTPTTYKAKQYLSYETLYANGSRTXQVGIYILSSSGDGAXXRDSSG 120
DB VSSSCGPGCHKGTPLTPTTYKAKQYLSYETLYANGSRTXQVGIYILSSSGDGAXXRDSSG 119

QY 121 SGKSRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEXHVLTAACHIDG 180
DB TGRSRRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEXHVLTAACHIDG 179

QY 181 KTYVKGTKLVRGFLKPKFKDGGRGANDSTSAPEQMKFQWIRVKRTHVPKGIKNAND 240
DB KTYVKGTKLVRGFLKPKFKDGGRGANDSTSAPEQMKFQWIRVKRTHVPKGIKNAND 239

QY 241 IGMDDYVALLLEKKPHKPKMKIGVSPPAKQLPGRHIFSGYNDPRLVTRFCDVKDE 300
DB IGMDDYVALLLEKKPHKPKMKIGVSPPAKQLPGRHIFSGYNDPRLVTRFCDVKDE 299

QY 301 TYDLLVQCDQAQPGASGYVYVWRKQKQKWKRIIGIFSGHGWDMNGSPDENVAVR 360
DB TYDLLVQCDQAQPGASGYVYVWRKQKQKWKRIIGIFSGHGWDMNGSPDENVAVR 359

QY 361 ITELKVAQICYWKGNLYDCREG 383
DB ITELKVAQICYWKGNLYDCREG 382

RESULT 4
Q8BZS4 PRELIMINARY; PRT; 382 AA.
AC Q8BZS4;
DT 01-MAR-2003 (TEMBLrel. 23, Created)
DT 01-MAR-2003 (TEMBLrel. 23, Last sequence update)
DE Mus musculus adult male cecum cDNA, RIKEN full-length enriched
DE library, clone:9130215B18 product:SERINE PROTEASE (HYPOTHETICAL 43.0
DE kDa PROTEIN) (PROTEASE, SERINE, 23) homolog.
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1] SEQUENCE FROM N.A.
RP STRAIN=C57BL/6J; TISSUE=Cecum;
RC MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
RX Carninci P., Hayashizaki Y.;
RT "High-efficiency full-length cDNA cloning.";
RN Meth. Enzymol. 303:19-44(1999).
[2]
RN SEQUENCE FROM N.A.
RP STRAIN=C57BL/6J; TISSUE=Cecum;
RC MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;
RX RIKEN FANTOM Consortium;
RT "Functional annotation of a full-length mouse cDNA collection.";
RN Nature 409:685-690(2001).
[3]
RN SEQUENCE FROM N.A.
RP STRAIN=C57BL/6J; TISSUE=Cecum;
RC The FANTOM Consortium;
RA the RIKEN Genome Exploration Research Group Phase I & II Team;
RT "Analysis of the mouse transcriptome based on functional annotation of
RT 60,770 full-length cDNAs.";
RN Nature 420:563-573(2002).
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[4] SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Cecum;
RX MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;
RA Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K., Itoh M.,
RA Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;
RT "Normalization and subtraction of cap-trapper-selected cDNAs to
RL Genome Res. 10:1617-1630(2000).
[5]
RN SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Cecum;
RX MEDLINE=20330913; PubMed=11076861; DOI=10.1101/gr.152600;
RA Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P.,
RA Konno H., Akiyama J., Nishi K., Kitsumai T., Tashiro H., Itoh M.,
RA Sumi N., Ishii Y., Nakamura S., Hazama S., Nishine T., Harada A.,
RA Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,
RA Fujiwaka S., Inoue K., Togawa Y., Izawa M., Ohara E., Watahiki M.,
RA Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuda S., Kawai J.,
RA Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.;
RT "RIKEN integrated sequence analysis (RISA) system-384-format
RL sequencing pipeline with 384 multicapillary sequencer.";
RN Genome Res. 10:1757-1771(2000).
[6]
RN SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Cecum;
RX Adachi J., Aizawa K., Akimura T., Arakawa T., Bono H., Carninci P.,
RA Fukushida S., Furuno M., Hanagaki T., Hara A., Hashizume W.,
RA Hayashida K., Hayatsu N., Hiramoto K., Hiraoka T., Hirozane T.,
RA Hori F., Inotani K., Ishii Y., Itoh M., Kagawa I., Kasukawa T.,
RA Kurihara C., Matsuyama T., Miyazaki A., Murata M., Nakamura M.,
RA Nishi K., Nomura K., Numazaki R., Ohno M., Ohsato N., Okazaki Y.,
RA Saito R., Saitoh H., Sakai C., Sakai K., Sakazume N., Sano H.,
RA Sasaki D., Shibata K., Shinagawa A., Shiraki T., Sogabe Y., Tagami M.,
RA Tagawa A., Takahashi F., Takaku-Akai H., Tanaka T., Tanaka T.,
RA Tomaru A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y.;
RL Submitted (JUL-2001) to the EMBL/GenBank/DBJ databases.
CC -!- SIMILARITY: Belongs to peptidase family S1.
DR EMBL; AK033671; BAC28420.1; -.
DR GO; GO:0004263; F:chymotrypsin activity; IEA.
DR GO; GO:0008233; F:peptidase activity; IEA.
DR GO; GO:0004299; F:trypsin activity; IEA.
DR GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR InterPro; IPR001254; Peptidase S1.
DR InterPro; IPR001314; Peptidase S1A.
DR InterPro; IPR009003; Pept Ser Cys.
DR PRINTS; PR00722; CHYMOTRYPSIN.
DR SMART; SM00020; tryp_Spc; 1.
DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
KW Hydrolase; Hypothetical protein; Protease; Serine protease.
SQ SEQUENCE 382 AA; 43147 MW; 556789818E12A081 CRC64;

Query Match 88.1%; Score 1860.5; DB 2; Length 382;
Best Local Similarity 89.6%; Pred. No. 1.7e-156;
Matches 343; Conservative 14; Mismatches 25; Indels 1; Gaps 1;

QY 1 MAGIPGLFLLFLLCAVGQVSPYAPWKPPTWPAVRLPVVLPQSTLNLAQDFGABAKLE 60
DB 1 MAGIPG-LFILLVLLCVFMQVSPYTPWKPPTWPAVRLPVVLPQSTLNLAQDFGABAKLE 59

QY 61 VSSSCGPGCHKGTPLTPTTYKAKQYLSYETLYANGSRTXQVGIYILSSSGDGAXXRDSSG 120
DB VSSSCGPGCHKGTPLTPTTYKAKQYLSYETLYANGSRTXQVGIYILSSSGDGAXXRDSSG 119

QY 121 SGKSRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEXHVLTAACHIDG 180
DB TGRSRRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEXHVLTAACHIDG 179

QY 181 KTYVKGTKLVRGFLKPKFKDGGRGANDSTSAPEQMKFQWIRVKRTHVPKGIKNAND 240
DB KTYVKGTKLVRGFLKPKFKDGGRGANDSTSAPEQMKFQWIRVKRTHVPKGIKNAND 239
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Db 298 TIKKPGGHIIFSGFNDRADQLYRFCSYDSNSDNLQYCDAESGSGTGGVYLRLKDP 357
 Qy 328 QOQWERKIIGIFSQHWDMNGSPQFNVAVRITPLKYAQICVWIKGNVLDREG 383
 Db 358 DKKNWKRKLIIAVSGHQWVGHGQKQYNNVAVRITPLKYAQICLWTHGNDANCAYG 413

RESULT 6 .
 Q8N3Z0 PRELIMINARY; PRT; 413 AA.
 AC Q8N3Z0;
 DT 01-OCT-2002 (TrEMBLrel. 22, Created)
 DT 25-OCT-2002 (TrEMBLrel. 22, Last sequence update)
 DT 25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
 DE Hypothetical protein PRSS35 (ENML522).
 GN Name=PRSS35; ORFName=UNQ522;
 OS Homo sapiens (Human).
 OC Eukaryota; Metazoa; Chordata; Cranata; Vertebrata; Euteleostomi;
 OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 NCBI_TaxID=9606;
 RN [1]
 RN SEQUENCE FROM N.A.
 RC TISSUE=Brain;
 RP MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.2426038999;
 RX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
 RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
 RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
 RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Haie F.,
 RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
 RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
 RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullaby S.J.,
 RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
 RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
 RA Fahey J., Helton E., Kettman M., Madan A.C., Shevchenko Y., Bouffard G.G.,
 RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
 RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
 RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
 RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
 RA Jones S.J., Marra M.A.;
 RT "Generation and initial analysis of more than 15,000 full-length human
 RT and mouse cDNA sequences";
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
 RN [2]
 RN SEQUENCE FROM N.A.
 RC TISSUE=Brain;
 RA Strausberg R.;
 RL Submitted (SEP-2002) to the EMBL/GenBank/DBJ databases.
 RN [3]
 RN SEQUENCE FROM N.A.
 RX MEDLINE=22887296; PubMed=12975309; DOI=10.1101/gr.1293003;
 RA Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,
 RA Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,
 RA Eaton D., Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S.,
 RA Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,
 RA Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,
 RA Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,
 RA Vanden R., Watanabe C., Winstead D., Woods K., Xie M.H., Yansura D.,
 RA Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,
 RA Godowski P.;
 RT "The secreted protein discovery initiative (SPDI), a large-scale
 RT effort to identify novel human secreted and transmembrane proteins: a
 RT bioinformatics assessment";
 RT Genome Res. 13:2265-2270(2003).
 CC -I- SIMILARITY: Belongs to peptidase family S1.
 DR EMBL, BC037170; AAH37170.1; -
 DR EMBL; AY358661; AAQ89024.1; -
 DR GO; GO:0004263; F:chymotrypsin activity; IEA.
 DR GO; GO:0008233; F:peptidase activity; IEA.
 DR GO; GO:0004295; F:trypsin activity; IEA.
 DR GO; GO:0006508; P:proteolysis and peptidolysis; IEA.


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RL Submitted (JUN-2004) to the EMBL/GenBank/DBJ databases.
CC -I- SIMILARITY: Belongs to peptidase family S1.
DR EMBL; AK031411; BAC27392.1; -
DR EMBL; BC075675; AA75675.1; -
DR MGD; MGI:2444800; P8835.
DR GO; GO:0005615; C:extracellular space; TAS.
DR InterPro; IPR001254; Peptidase_S1.
DR InterPro; IPR001314; Peptidase_S1A.
DR InterPro; IPR009003; Pept_Ser_Cys.
DR Pfam; PF00089; Trypsin; 1.
DR PRINTS; PR00722; CHYMOTRYPSIN.
DR SMART; SM00020; Tryp_SPC; 1.
DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR HydroLase; Protease; Serine protease.
KW SEQUENCE 409 AA; 45787 MW; 5E22D4A908E7EF5 CRC64;

Query Match 47.5%; Score 1003.5; DB 2; Length 409;
Best Local Similarity 48.6%; Pred. No. 2.1e-80;
Matches 191; Conservative 68; Mismatches 85; Indels 47; Gaps 5;

QY 31 TWPAYRLPVLPVLPSTLNLAKEPFGAEAKLEVSSCGPQCHKGTPLPYKEAKQYLSYETL 90
DB 28 TWHLSRIPQVVSNTIHLASPTFQADAGVVVKATVCGIECQELPAPSLQLESLSYETI 87

QY 91 YANGSRTEKXQVGIYL-----SSSGDGAAXRDSGSGKRRKQIYGYDSRFSIFGKDF 144
DB 88 FENGTRITLVKQVGLVLETRNSVKG-----HPRRRQVGTDSRFSILDKRF 138

QY 145 LLNYPFSTSVKLTGCTGTLVAEXHVLTAACHDGNKYVKGQKLRVGLKPKFKDG-- 202
DB 139 ATNFPNTAVKLTGCGTGLVSPNHVLTAAHCVHDGKDYVKGSKLRVGLKVRNKGGRK 198

QY 203 -GRGA-----NDSTSAMPEQM-----KQWIRVGRTHVPKG 232
DB 199 KRGSKRSRREAESAGSQAHLESTTQRPCKSRGRPRVTQGRPSQWTRVKSHTPKG 258

QY 233 WIKGNANDIGMDYDYLLELKKPKKFMKIGVSPPAKQLPGRIHFGSDVNDRPNGLVY 292
DB 259 WVGENGGLALDYDYLLELKKRAHKQHWELGVSPITKLPGRRIHFGSDNDRDQLVY 318

QY 293 RFDVDRDEYDLYQQCDQAQGSAGSGVYVVMKQKQKWERKIIGFSGHQWDMNGSP 352
DB 319 RFCSVSEESNDLLYQCDABAGSTGSGIYLRLEKPEQKWKRIYAVYSGHQWVDHGQV 378

QY 353 QDPNVAVRITPLKYAQICVYKIGNYLDREG 383
DB 379 KDYNVAVRITPLKYAQICLWIHGNAANCAYG 409

RESULT 8
Q8COL5 PRELIMINARY; PRT; 409 AA.
AC Q8COL5;
DT 01-MAR-2003 (TrEMBLrel. 23, Created)
DT 01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE Mus musculus 6 days neonate head cDNA, RIKEN full-length enriched
DE library, clone:5430417J04 product:similar to DJ223E3.1 (PUTATIVE
DE SECRETED PROTEIN ZS1G13), full insert sequence.
GN Names=Pras35;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RN SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Head;
RX MEDLINE=99279233; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
RA Carninci P., Hayashizaki Y.;
RT "High-efficiency full-length cDNA cloning.";
RL Meth. Enzymol. 303:19-44(1999).
RN [2]
RN SEQUENCE FROM N.A.

RC STRAIN=C57BL/6J; TISSUE=Head;
RX MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;
RA RIKEN FANTOM Consortium;
RT "Functional annotation of a full-length mouse cDNA collection.";
RL Nature 409:685-690(2001).
RN [3]
RN SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Head;
RA the RIKEN Genome Exploration Research Group Phase I & II Team;
RT "Analysis of the mouse transcriptome based on functional annotation of
RT 60,770 full-length cDNAs.";
RL Nature 420:563-573(2002).
RN [4]
RN SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Head;
RX MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;
RA Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K., Itoh M.,
RA Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;
RT "Normalization and subtraction of cap-trapper-selected cDNAs to
RT prepare full-length cDNA libraries for rapid discovery of new genes.";
RL Genome Res. 10:1617-1630(2000).
RN [5]
RN SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Head;
RX MEDLINE=1076861; DOI=10.1101/gr.152600;
RA Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P.,
RA Konno H., Akiyama Y., Nishi K., Kitsunai T., Tashiro H., Itoh M.,
RA Sumi N., Ishii Y., Nakamura S., Hazama M., Nishine T., Harada A.,
RA Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,
RA Fujiwake S., Inoue K., Togawa Y., Izawa M., Ohara E., Watahiki M.,
RA Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,
RA Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.;
RT "RIKEN integrated sequence analysis (RISA) system-384-format
RT sequencing pipeline with 384 multicapillary sequencer.";
RL Genome Res. 10:1757-1771(2000).
RN [6]
RN SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Head;
RA Adachi J., Aizawa K., Akimura T., Arakawa T., Bono H., Carninci P.,
RA Fukuda S., Furuno M., Hanagaki T., Hara A., Hashizume W.,
RA Hayashida K., Hayatsu N., Hiramoto K., Hiraoka T., Hirozane T.,
RA Hori F., Inotani K., Ishii Y., Itoh M., Kagawa I., Kasukawa T.,
RA Katoh H., Kawai J., Kojima Y., Kondo S., Konno H., Kouda M., Koya S.,
RA Kurihara C., Matsuyama T., Miyazaki A., Murata M., Nakamura M.,
RA Nishi K., Nomura K., Numazaki R., Ohno M., Ohsato N., Okazaki Y.,
RA Saito R., Saitoh H., Sakai C., Sakai K., Sakazume N., Sano H.,
RA Sasaki D., Shibata K., Shinagawa A., Shiraki T., Sogabe Y., Tagami M.,
RA Tagawa A., Takahashi F., Takaku-Akahira S., Takeda Y., Tanaka T.,
RA Tomaru A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y.;
RL Submitted (JUL-2001) to the EMBL/GenBank/DBJ databases.
CC -I- SIMILARITY: Belongs to peptidase family S1.
DR EMBL; AK030671; BAC27073.1; -
DR MGD; MGI:2444800; P8835.
DR GO; GO:0005615; C:extracellular space; TAS.
DR InterPro; IPR001254; Peptidase_S1.
DR InterPro; IPR001314; Peptidase_S1A.
DR InterPro; IPR009003; Pept_Ser_Cys.
DR Pfam; PF00089; Trypsin; 1.
DR PRINTS; PR00722; CHYMOTRYPSIN.
DR SMART; SM00020; Tryp_SPC; 1.
DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
KW HydroLase; Protease; Serine protease.
KW SEQUENCE 409 AA; 45847 MW; ASAC1C582BFD352E CRC64;

Query Match 47.3%; Score 998.5; DB 2; Length 409;
Best Local Similarity 48.6%; Pred. No. 5.8e-80;
Matches 190; Conservative 68; Mismatches 86; Indels 47; Gaps 5;

QY 31 TWPAYRLPVLPVLPSTLNLAKEPFGAEAKLEVSSCGPQCHKGTPLPYKEAKQYLSYETL 90
DB 28 TWHLSRIPQVVSNTIHLASPTFQADAGVVVKATVCGIECQELPAPSLQLESLSYETI 87
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QY 91 YANGSRTEXOVGIYIL-----SSSGDAXXRDSSGSGKRRKQIYGYVDSRFSIFGKDF 144
Db 88 FENGTRTLVQVGLLEPTRNSVKG-----HPRRRQVYGTDSRFSILDKRF 138
QY 145 LLNVPRSTSVKLGCTGLVAEXHVLTAACHIDGKTVYKQKLRVGLPKPKFKDG-- 202
Db 139 ATNFPFNIAVLSTGCGTLVSPNHLVTAACHVDGKYVKGSKLRVGLKVRNKGGRK 198
QY 203 -GRGA-----NDSTSAMPEQM-----KFQMRVVRKTHVPKG 232
Db 199 KRGSKRSREAESAGSOAHLRESITQRPKKSRGPRVYQGPSPQWTRVKSITHPKG 258
QY 233 WIKGNANDIGMDYDVALLELKKPKHKFKMIGVSPPAKQLPGRIHFGSDNDRPNLVY 292
Db 259 WVRGSGSLADYDVALLELKRHKQHMELGVSPTITKLPGRIHFGSDNDRDQLVY 318
QY 293 RFDVNDKTDLYAQCDAPGASGVYVVRMKRQOQKWKRIIGFSGHWDVNGSP 352
Db 319 RFGVSSESNLLYQYCDAGSTGSGIYLRKPKQKWKRIYAVYSGHWDVNGVQ 378
QY 353 QDNVAVRITPLKYAQICYWIKGNIDCDREG 383
Db 379 KDYNVAVRITPLKYAQICLWIHGNAACVAG 409

RESULT 9
Q6GML6 PRELIMINARY; PRT; 418 AA.
ID Q6GML6
AC Q8CODE PRELIMINARY; PRT; 409 AA.
DT 05-JUL-2004 (TREMBLrel. 27, Created)
DT 05-JUL-2004 (TREMBLrel. 27, Last sequence update)
DE zgc:91804.
GN ORFNames=zgc:91804;
OS Brachydanio rerio (Zebrafish) (Danio rerio).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
OC Cyprinidae; Danio.
OX NCBI_TaxID=7955;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=whole;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Scapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Uedin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahney J., Heiton E., Ketterman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakeley R.W., Touchman J.W., Green E.D., Dickinson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalak U., Smallos D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RA "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]
RP SEQUENCE FROM N.A.
RC TISSUE=whole;
RX Strausberg R.;
RA Submitted (JUN-2004) to the EMBL/GenBank/DBJ databases.
DR ENBL; BC074028; AAH74028.1; -.
DR ZFIN; ZDB-GENE-040704-55; zgc:91804.
DR GO; GO:0008233; F:peptidase activity; IEA.
DR GO; GO:0004295; F:trypsin activity; IEA.
DR GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
```

```
DR InterPro; IPR001254; Peptidase_S1.
DR InterPro; IPR009003; Pept_Ser_Cys.
DR Pfam; PF00089; Trypsin; 1.
DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
KW Hydrolase; Protease; Serine protease.
SQ SEQUENCE 418 AA; 47270 MW; AF53345CEB94F720 CRC64;

Query Match 47.1%; Score 994.5; DB 2; Length 418;
Best Local Similarity 45.8%; Pred. No. 1.3e-79;
Matches 193; Conservative 63; Mismatches 122; Indels 43; Gaps 2;

QY 1 MAGIFGLLFLFLCAVQVSPYAPWKPWTWPAYRLPVVLQSTLNLAQPFQGAQKLE 60
Db 1 MGPEVPLTLLLSALAVLGSTTVDPTGDTWTPQRIPLVQEKQTVHLESSEFLAKPQND 60
QY 61 VSSSGCPQCHKGTPLPTVKEAQVLSYETLYANGSRTEXQVGIYILSSSGDCAXXRDSGS 120
Db 61 LHGICIGEQQLRPFSLDLEQLLSYETWYDNGTRTLTTVTVDLNVNDWT-----GAS 116
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLGSTGCTGLVAEXHVLTAACHIDG 180
Db 117 QLHTRHREVYGTDRFTITDKQYSMKYPFSTSVKISTGCGVLVSPKHVLTAAACHIHG 176
QY 181 KTYVGTOKLRVGLKP-----KFKD 201
Db 177 TDYLDGVQKLSVGLKERSRRKNGRKGKRGKQKHHEEVEVDENGEIVEXQERKSKG 236
QY 202 GGRGANDSTSAMPEOMKQWIRVKTVPKGIKGNANDIGMDYDVALLELKKPKHKRF 261
Db 237 KGRNRSRSTDSQSPFRWTRVKQVQKGFKGISENLADYDVALLELRAQTKRF 296
QY 262 KIGVPPAKQLPGRHFGSDNDRPGLNLYRVCDFDKDDEYDLYQQCDAPGASGVY 321
Db 297 DLGVIPSVKCLPAGRIHFGSDNDRPGLNLYRVCDFDKDDEYDLYQQCDAPGASGVY 356
QY 322 VRMVTRQOQWKRIIGFSGHWDVNGSPDFNVAVRITPLKYAQICYWIKGNVLDRC 381
Db 357 IRLKEPFGKKKWKRIIGVFGSHQWVDVNGQDYNVAVRITPLKYAQICRWVHGSSQCR 416
QY 382 E 382
Db 417 D 417

RESULT 10
Q8CODE PRELIMINARY; PRT; 409 AA.
ID Q8CODE PRELIMINARY; PRT; 409 AA.
AC Q8CODE; PRELIMINARY; PRT; 409 AA.
DT 01-MAR-2003 (TREMBLrel. 23, Created)
DT 01-MAR-2003 (TREMBLrel. 23, Last sequence update)
DT 01-MAR-2004 (TREMBLrel. 26, Last annotation update)
DE Mus musculus 13 days embryo male testis cDNA, RIKEN full-length
DE enriched library clone:603046M24 product:similar to DJ223E3.1
DE (PUTATIVE SECRETED PROTEIN ZSGL3), full insert sequence.
GN Name=Prss35;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Testis;
RX MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
RA Carninci P., Hayashizaki Y.;
RT "High-efficiency full-length cDNA cloning.";
RL Meth. Enzymol. 303:19-44(1999).
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Testis;
RX MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;
RA RIKEN PANTOM Consortium;
RT "Functional annotation of a full-length mouse cDNA collection.";
RL Nature 409:685-690(2001).
```

RN [3]
 RP SEQUENCE FROM N.A.
 RC STRAIN=C57BL/6J; TISSUE=Testis;
 RA The FANTOM Consortium;
 RA the RIKEN Genome Exploration Research Group Phase I & II Team;
 RT "Analysis of the mouse transcriptome based on functional annotation of
 RT 60,770 full-length cDNAs";
 RL Nature 420:563-573(2002).
 RN [4]
 RP SEQUENCE FROM N.A.
 RC STRAIN=C57BL/6J; TISSUE=Testis;
 RX MEDLINE=20493374; PubMed=11042159; DOI=10.1101/gr.145100;
 RA Carninci P., Shibata Y., Hayatsu M., Sugahara Y., Shibata K., Itoh M.,
 RA Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.,
 RT "Normalization and subtraction of cap-trapper-selected cDNAs to
 RT prepare full-length cDNA libraries for rapid discovery of new genes";
 RL Genome Res. 10:1617-1630(2000).
 RN [5]
 RP SEQUENCE FROM N.A.
 RC STRAIN=C57BL/6J; TISSUE=Testis;
 RX MEDLINE=20530913; PubMed=11076861; DOI=10.1101/gr.152600;
 RA Shibata K., Itoh M., Aizawa K., Nagao S., Sasaki N., Carninci P.,
 RA Konno H., Akiyama J., Nishi K., Kiteunai T., Tashiro H., Itoh M.,
 RA Sumi N., Ishii Y., Nakamura S., Hazama M., Nishine T., Harada A.,
 RA Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,
 RA Fujiwaka S., Inoue K., Togawa Y., Izawa M., Ohara E., Watahiki M.,
 RA Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,
 RA Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.,
 RT "RIKEN integrated sequence analysis (RISA) system-384-format
 RT sequencing pipeline with 384 multicapillary sequencer";
 RL Genome Res. 10:1757-1771(2000).
 RN [6]
 RP SEQUENCE FROM N.A.
 RC STRAIN=C57BL/6J; TISSUE=Testis;
 RA Adachi J., Aizawa K., Akimura T., Arakawa T., Bono H., Carninci P.,
 RA Fukuda S., Furuno M., Hanagaki T., Hara A., Hashizume W.,
 RA Hayashida K., Hayatsu M., Hiramoto K., Hiraoka T., Hirozane T.,
 RA Hori F., Imotani K., Ishii Y., Itoh M., Kagawa I., Kasukawa T.,
 RA Katoh H., Kawai J., Kojima Y., Kondo S., Konno H., Kouda M., Koya S.,
 RA Kurihara C., Matsuyama T., Miyazaki A., Murata M., Nakamura M.,
 RA Nishi K., Nomura K., Numazaki R., Ohno M., Ohsato N., Okazaki Y.,
 RA Saito R., Saitoh H., Sakai C., Sakai K., Sakazume N., Sano H.,
 RA Sasaki D., Shibata K., Shingawa A., Shiraki T., Sogabe Y., Tagami M.,
 RA Tagawa A., Takahashi F., Takaku-Akashira S., Takeda Y., Tanaka T.,
 RA Tomaru A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y.,
 RL Submitted (JUL-2001) to the EMBL/GenBank/DBJ databases.
 DR EMBL; AK031644; BAC27491.1; -;
 DR MGD; MGI:2444800; Pss635.
 DR GO; GO:0005615; C:extracellular space; TAS.
 DR InterPro; IPR001254; Peptidase_S1.
 DR InterPro; IPR009003; Pept_Ser_Cys.
 DR Pfam; PF00089; Trypsin; 1.
 DR SMART; SM00020; Tryp_SPC; 1.
 KW Hydrolyase; Protease; Serine protease.
 SQ SEQUENCE 409 AA; 45736 MW; B9C9B91E6800719C CRC64;
 Query Match 47.0%; Score 992.5; DB 2; Length 409;
 Best Local Similarity 48.3%; Pred. No. 2e-79;
 Matches 189; Conservative 70; Mismatches 85; Indels 47; Gaps 5;
 QY 31 TWPAYLPVVLPOSTLNLAKPDFGAELKLEVVSSCGPQCHGTPPLTYKEAKQYLSVETL 90
 DB 28 TWHLRIPQVSVSTLHLSPTQADQVVKATVCIGECQELPAPLSQLESLSYETI 87
 QY 91 YANGSRTEQVGIYIL-----SSSDGAXXRDSGSGSKRRKQIYGVDSRFSIFGKDF 144
 DB 88 FENGTRTLRVKQGLVLETRNSVKGK-----HPRRRQVYGTDSRFSILDKPF 138
 QY 145 LNNYPRSTSVKLTGCTGTVAEXHVLTAACHDGTYYVKGQKLRVGLPKFPGKDG-- 202
 DB 139 ATNFPNTAVKLTSGCGSLVSPNHLTAANCVDHGDYVKGSKLRLVGLKMRNGGRK 198
 QY 203 -GRGA-----NDSTSAMPEQM-----XEQWIRVKRTHVPGK 232

DB 199 KRGSKRREAREASQSOAHLRESITTPQPKSRGRPRVTQGRPSFQWTRVASTHPKG 258
 QY 233 WTKGNANDIGMDYVYALLELKPHKFKFKMKGIVSPPAKQLPGRIHPSGVNDNRPNLVY 292
 DB 259 WVRGEGGLADYVYALLELKRAHKQHMELGVSPTITKLPQGGIHFSGFDNRDEQLVY 318
 QY 293 RFGCDVXDYDLYLQOCDAOPGASGVVYVWKKROQKWKIIGIFSGHGVNDKNSP 352
 DB 319 RFGVSEESNDLYQYCDABAGSTGSIYLRLEKPGQKWKRIYAVISGHQWVDVHGQV 378
 QY 353 QDFNVAVRITPLKYAQICWIKNGYLDCREG 383
 DB 379 KDYNVAVRITPLKYAQICLWIGHNANCAVG 409
 RESULT 11
 GSEP_BACLI
 ID GSEP_BACLI STANDARD; PRT; 316 AA.
 AC P80057;
 DT 01-MAY-1992 (Rel. 22, Created)
 DT 01-OCT-1993 (Rel. 27, Last sequence update)
 DT 25-OCT-2004 (Rel. 45, Last annotation update)
 DE Glutamyl endopeptidase precursor (EC 3.4.21.19) (Glutamate specific
 endopeptidase) (GSE).
 GN Name:blase;
 OS Bacillus licheniformis.
 OC Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus.
 OX NCBI_TaxID=1402;
 RN [1]
 RP SEQUENCE FROM N.A., AND PARTIAL SEQUENCE.
 RC STRAIN=ATCC 14580;
 RX MEDLINE=93054737; PubMed=1429718;
 RA Kakudo S., Kikuchi N., Kitadokoro K., Fujiwara T., Nakamura E.,
 RA Okamoto H., Shin M., Tanaki M., Teraoka H., Tsuzuki H., Yoshida N.;
 RT "Purification, characterization, cloning, and expression of a glutamic
 RT acid-specific protease from Bacillus licheniformis ATCC 14580";
 RJ J. Biol. Chem. 267:23782-23788(1992).
 RN [2]
 RP SEQUENCE OF 95-316.
 RX MEDLINE=92155199; PubMed=1346764;
 RA Svendsen I., Bredsdal K.;
 RT "Isolation and amino acid sequence of a glutamic acid specific
 RT endopeptidase from Bacillus licheniformis";
 RJ Eur. J. Biochem. 204:165-171(1992).
 CC -!- FUNCTION: Specific for hydrolysis of peptide bonds on the carboxyl
 CC side of acidic amino acid residues, with a strong preference for
 CC Glu.
 CC -!- CATALYTIC ACTIVITY: Preferential cleavage: Asp|-Xaa, Glu|-Xaa.
 CC -!- SUBCELLULAR LOCATION: Secreted.
 CC -!- SIMILARITY: Belongs to the peptidase S1B family.
 CC -----
 CC This SWISS-PROT entry is copyright. It is produced through a collaboration
 CC between the Swiss Institute of Bioinformatics and the EMBL outstation -
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 CC or send an email to license@sib-sib.ch).
 CC -----
 CC EMBL; D10060; BAA00949.1; -;
 DR FIR; A45134; A45134.
 DR MEROPS; S01.271; -;
 DR InterPro; IPR000126; Pept_S1B_AS.
 DR InterPro; IPR009003; Pept_Ser_Cys.
 DR InterPro; IPR001254; Peptidase_S1.
 DR InterPro; IPR008256; Peptidase_S1B.
 DR Pfam; PF00089; Trypsin; 1.
 DR PRINTS; PR00839; V8PROTEASE.
 DR SMART; SM00020; Tryp_SPC; 1.
 DR PROSITE; PS00672; V8_HIS; 1.
 DR PROSITE; PS00673; V8_SER; 1.
 KW Direct protein sequencing; Hydrolase; Serine protease; Signal.

[illegible][illegible]


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169 YGCSATLIGPRTVLTAAHCLYSHEDKDWL---SBYLPV-----PGLNGSTA--- 212
215 EQMKFQIRVKRTHVPKGTWKGNDIG--MDYDVALLELKKPHKRFKMGKIVSPPAKOL 272
213 DDAPFGAFTYESAVLQGFIDNYQGYGYSVIPWDLGIITLKQDVGTNLGWLGYA-NYDDL 271
273 PGCRIHPSGYDNDRP-GNLVYRFCDVKDETDLLYQQ--CDAQPCASGVGVYVRMMKROQ 329
272 GDFANLVGYPGDGPMGTMMKASCEVHAENIAPFYQYDCDTFFGSSSSSVYAYDTKSKQ 331
330 QKWERKIIGFSGHQWDMNGSPQDFNVAVRITPLKYAICYWIKGNY 377
332 R-----IITG---VNVAESP-DANTAVRLN---AANVQWINSLY 363

RESULT 15
Q987W6
ID Q987W6 PRELIMINARY; PRT; 271 AA.
AC Q987W6;
DT 01-OCT-2001 (TrEMBLrel. 18, Created)
DT 01-OCT-2001 (TrEMBLrel. 18, Last sequence update)
DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE Glutamic acid specific endonopeptidase.

```

OS Rhizobium loti (Mesorhizobium loti).
OC Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
OC Phyllobacteriaceae; Mesorhizobium.
OX NCBI_TaxID=381;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=MAFF303099;
RX MEDLINE=21082930; PubMed=11214968;
RA Kaneko T., Nakamura Y., Sato S., Asamizu E., Kato T., Sasamoto S.,
RA Watanabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura T.,
RA Kishida Y., Kiyokawa C., Kohara M., Matsumoto M., Matsuno A.,
RA Mochizuki Y., Nakayama S., Nakayama K., Nakazaki N., Shimpo S., Sugimoto M.,
RA Takeuchi C., Yamada M., Tabata S.;
RT "Complete genome structure of the nitrogen-fixing symbiotic bacterium
RT Mesorhizobium loti.";
RL DNA Ref. 7:311-338(2000).
CC -1- SIMILARITY: Belongs to peptidase family S1.
DR EMBL, AP003010; BAB5084.1; -.
DR GO; GO:0004263; F:chymotrypsin activity; IEA.
DR GO; GO:0004295; F:trypsin activity; IEA.
DR GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR InterPro; IPR001254; Peptidase_S1.
DR InterPro; IPR001314; Peptidase_S1A.
DR InterPro; IPR008256; Peptidase_S1B.
DR InterPro; IPR009003; Pept_Ser_Cys.
DR Pfam; PF00089; Trypsin; 1.
DR PRINTS; PR00722; CHYMOTRYPSIN.
DR PRINTS; PR00839; VAPROTEASE.
DR SMART; SM00020; Tryp_SPC; 1.
DR PROSITE; PS00240; TRYPSIN_DOM; 1.
DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN 1.
KW Complete proteome; Hydrolase; Protease; Serine protease.
SQ SEQUENCE 271 AA; 30385 MW; 37659307C63D0D9C CRC64;
Query Match 7.4%; Score 156; DB 2; Length 271;
Best Local Similarity 29.0%; Pred. No. 2e-05;
Matches 67; Conservative 21; Mismatches 91; Indels 52; Gaps 12

158	QY	TGCTGTLTVAEXHVLTTAAHCHIDGKTYVKGFKLURVGF-LKP	KFKDGGGANDSTSAMPEQ	211
159	QY			
160	QY			
161	QY			
162	QY			
163	QY			
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167	QY			
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214	QY			
215	QY			
216	QY			
217	QY	MKFQWLRVKRTHV-----PKGWIKNANDIGWDYDYLLEKPKH	KFKMKIGVSPPA	269
218	QY			
219	QY			
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256	QY			
257	QY			
258	QY			
259	QY			
260	QY			
261	QY			
262	QY			
263	QY			
264	QY			
265	QY			
266	QY			

Oy 324 MWKRQQQKWKIIGI-----FSGHQWYDMNGSPQD-----FNVAVRITP 363
Db 210 -WIHRQQAGPPVIVAVHTAGPRPHSGGAMGCRPGVPLAPAGLFNRCVRLTP 259

Search completed: July 1, 2005, 21:08:10
Job time : 96.7095 secs

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: July 1, 2005, 20:52:53 ; Search time 100.435 Seconds
(without alignments)
1509.530 Million cell updates/sec

Title: US-09-658-677-15
Perfect score: 2131
Sequence: 1 MAGIPGLFLFLFLCAVQ.....IKGNVLDRCRGTVFPFGSN 392

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2105692 seqs, 386760381 residues

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : A_Geneseq_16Dec04:*
1: Geneseq1980s:*
2: Geneseq1990s:*
3: Geneseq2000s:*
4: Geneseq2001s:*
5: Geneseq2002s:*
6: Geneseq2003as:*
7: Geneseq2003bs:*
8: Geneseq2004s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Match	Query Length	ID	Description
1	2131	100.0	392	4	AAB48973 Human Zsi
2	2085	97.8	392	4	AAB48972 Human Zsi
3	2080	97.6	383	2	AAY08660 WO9927094
4	2080	97.6	383	2	AAY08657 Human tra
5	2080	97.6	383	2	AAY13390 Amino aci
6	2080	97.6	383	3	AAY88277 Human TAN
7	2080	97.6	383	3	AAY87270 Human sig
8	2080	97.6	383	3	AAY53627 A bone ma
9	2080	97.6	383	3	AAB25618 Protein e
10	2080	97.6	383	3	AAB25592 Protein e
11	2080	97.6	383	3	ADC78573 Human PRO
12	2080	97.6	383	4	AAB80258 Human PRO
13	2080	97.6	383	4	AAB48974 Human Zsi
14	2080	97.6	383	4	AAB29048 Human PRO
15	2080	97.6	383	6	ABU58424 Human PRO
16	2080	97.6	383	6	ABU71636 Human PRO
17	2080	97.6	383	6	ABU87972 Novel hum
18	2080	97.6	383	6	ABU84287 Human sec
19	2080	97.6	383	6	ABR66161 Human sec
20	2080	97.6	383	6	ABR65551 Human sec
21	2080	97.6	383	6	ABU99491 Human sec
22	2080	97.6	383	6	ABU82730 Human PRO
23	2080	97.6	383	6	ABU89851 Novel hum
24	2080	97.6	383	6	ABU71491 Human PRO
25	2080	97.6	383	6	ABR68100 Human sec

26	2080	97.6	383	6	ABU96153	Novel hum
27	2080	97.6	383	6	ABU92584	Human sec
28	2080	97.6	383	6	ABO08661	Human sec
29	2080	97.6	383	6	ABO02713	Human sec
30	2080	97.6	383	6	ABR74867	Human sec
31	2080	97.6	383	6	ABR94629	Human sec
32	2080	97.6	383	6	ABU85602	Human PRO
33	2080	97.6	383	6	ABU98762	Novel hum
34	2080	97.6	383	6	ABU97977	Novel hum
35	2080	97.6	383	6	ABU91683	Novel hum
36	2080	97.6	383	6	ABU71937	Human sec
37	2080	97.6	383	6	ABU89376	Human PRO
38	2080	97.6	383	6	ABU86217	Human sec
39	2080	97.6	383	6	ABU67430	Human sec
40	2080	97.6	383	6	ABU80458	Human PRO
41	2080	97.6	383	6	ABO01820	Novel hum
42	2080	97.6	383	6	ABR99376	Human sec
43	2080	97.6	383	6	ABR98766	Human sec
44	2080	97.6	383	6	ABO16289	Human sec
45	2080	97.6	383	6	ABR92189	Human sec

ALIGNMENTS

RESULT 1
AAB48973
ID AAB48973 standard; protein; 392 AA.
XX
AC AAB48973;
XX
DT 27-MAR-2001 (first entry)
XX
DE Human Zsig13 variant #2, SEQ ID NO:15.
XX
KW Human Zsig13; serine protease; chromosome 11q22.1; elastase homologue;
KW glutamyl endopeptidase homologue; factor X homologue; trypsin homologue;
KW trypsinogen homologue; mast cell protease homologue;
KW collagenase homologue; protein degradation; food processing; brewing;
KW alcohol production; laundry detergent component.
XX
OS Homo sapiens.
XX
PN US6153420-A.
XX
PD 28-NOV-2000.
XX
PF 04-MAY-1998; 98US-00072384.
XX
PR 24-APR-1997; 97US-0044185P.
PR 17-APR-1998; 98US-00062142.
XX
PA (ZYMO) ZYMOGENETICS INC.
XX
PI Sheppard PO;
XX
DR WPI; 2001-060090/07.
DR N-PSDB; AAC91783.
XX
PT New isolated serine protease (designated Zsig13), useful in industrial processes to degrade unwanted proteins or alter the characteristics of protein-containing composition, as well as in industrial applications (e.g. brewing).
PT
PT
PT
PT
PS Claim 1; Col 35-38; 26pp; English.
XX
CC The invention relates to human Zsig13 proteins (AAB48972-B48974), and to DNA encoding them (AAC91782-C91784). The invention also relates to expression vectors and host cells comprising a human Zsig13 DNA, and the recombinant production of a human Zsig13 protein or its precursor. Zsig13 is a serine protease, and has significant homology to Bacillus licheniformis glutamyl endopeptidase, human clotting factor X, human elastase, rat mast cell protease, Streptomyces griseus trypsin, bovine

CC trypsinogen, and Hypoderma lineatum collagenase. The gene encoding human
CC Zsig13 is located on chromosome 11q22.1. Zsig13 is useful in industrial
CC processes to degrade unwanted proteins or alter the characteristics of
CC protein-containing compositions. It may also be used in industrial
CC applications in which proteases are utilised, including food processing,
CC brewing and alcohol production, and as a component of a laundry
CC detergent. The present sequence represents a human Zsig13 variant
XX
SQ Sequence 392 AA;

Query Match 100.0%; Score 2131; DB 4; Length 392;
Best Local Similarity 100.0%; Pred. No. 4.4e-151; Mismatches 0; Indels 0; Gaps 0;
Matches 392; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLCAVGVSPYSAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPGCHKTPLPTYEAKQYLSYETLYANGSRRTETQVGIYILSSSGDGAQHRS DGS 120
Db 61 VSSSCGPGCHKTPLPTYEAKQYLSYETLYANGSRRTETQVGIYILSSSGDGAQHRS DGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180

QY 181 KTYVKGTKLRVGLFKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240
Db 181 KTYVKGTKLRVGLFKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240

QY 241 IGMVDYALLESKPKHKKFKMKGIVSPPAKQLPGGRIHPSGYNDNRPGNLVTRFCVDKDE 300
Db 241 IGMVDYALLESKPKHKKFKMKGIVSPPAKQLPGGRIHPSGYNDNRPGNLVTRFCVDKDE 300

QY 301 TYDLLYQCCDAQPGASGSGVYVRMKRQKQKWERKIIGIFSGHQWVDMNGSPQDENVAVR 360
Db 301 TYDLLYQCCDAQPGASGSGVYVRMKRQKQKWERKIIGIFSGHQWVDMNGSPQDENVAVR 360

QY 361 ITPKLYAQICYWIKGNLYDCREGDVTFFPPGNS 392
Db 361 ITPKLYAQICYWIKGNLYDCREGDVTFFPPGNS 392

RESULT 2
AAB48972
ID AAB48972 standard; protein; 392 AA.
AC AAB48972;
XX
XX 27-MAR-2001 (first entry)
XX Human Zsig13 variant #1, SEQ ID NO:2.
XX
KW Human Zsig13; serine protease; chromosome 11q22.1; elastase homologue;
KW glutamyl endopeptidase homologue; factor x homologue; trypsin homologue;
KW trypsinogen homologue; mast cell protease homologue;
KW collagenase homologue; protein degradation; food processing; brewing;
KW alcohol production; laundry detergent component.
XX
OS Homo sapiens.
XX
XX US6153420-A.
XX
XX 28-NOV-2000.
XX
XX 04-MAY-1998; 98US-00072384.
XX
XX 24-APR-1997; 97US-0044185P.
XX 17-APR-1998; 98US-00062142.
XX
XX (ZYMO) ZYMOGENETICS INC.
XX
XX Sheppard PO;

XX
DR WPI; 2001-060090/07.
DR N-PSDB; AAC91782.
XX
PT New isolated serine protease (designated Zsig13), useful in industrial
PT processes to degrade unwanted proteins or alter the characteristics of
PT protein-containing composition, as well as in industrial applications
PT (e.g. brewing).
XX
XX Claim 1; Col 25-28; 26pp; English.
XX
XX The invention relates to human Zsig13 proteins (AAB48972-B48974), and to
CC DNA encoding them (AAC91782-C91784). The invention also relates to
CC expression vectors and host cells comprising a human Zsig13 DNA, and the
CC recombinant production of a human Zsig13 protein or its precursor. Zsig13
CC is a serine protease, and has significant homology to Bacillus
CC licheniformis glutamyl endopeptidase, human clotting factor X, human
CC elastase, rat mast cell protease, Streptomyces griseus trypsin, bovine
CC trypsinogen, and Hypoderma lineatum collagenase. The gene encoding human
CC Zsig13 is located on chromosome 11q22.1. Zsig13 is useful in industrial
CC processes to degrade unwanted proteins or alter the characteristics of
CC protein-containing compositions. It may also be used in industrial
CC applications in which proteases are utilised, including food processing,
CC brewing and alcohol production, and as a component of a laundry
CC detergent. The present sequence represents a human Zsig13 variant
XX
SQ Sequence 392 AA;

Query Match 97.8%; Score 2085; DB 4; Length 392;
Best Local Similarity 98.2%; Pred. No. 1.2e-147; Mismatches 6; Indels 0; Gaps 0;
Matches 385; Conservative 1; Mismatches 6; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLCAVGVSPYSAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPGCHKTPLPTYEAKQYLSYETLYANGSRRTETQVGIYILSSSGDGAQHRS DGS 120
Db 61 VSSSCGPGCHKTPLPTYEAKQYLSYETLYANGSRRTETQVGIYILSSSGDGAQHRS DGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180

QY 181 KTYVKGTKLRVGLFKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240
Db 181 KTYVKGTKLRVGLFKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240

QY 241 IGMVDYALLESKPKHKKFKMKGIVSPPAKQLPGGRIHPSGYNDNRPGNLVTRFCVDKDE 300
Db 241 IGMVDYALLESKPKHKKFKMKGIVSPPAKQLPGGRIHPSGYNDNRPGNLVTRFCVDKDE 300

QY 301 TYDLLYQCCDAQPGASGSGVYVRMKRQKQKWERKIIGIFSGHQWVDMNGSPQDENVAVR 360
Db 301 TYDLLYQCCDAQPGASGSGVYVRMKRQKQKWERKIIGIFSGHQWVDMNGSPQDENVAVR 360

QY 361 ITPKLYAQICYWIKGNLYDCREGDVTFFPPGNS 392
Db 361 ITPKLYAQICYWIKGNLYDCREGDVTFFPPGNS 392

RESULT 3
AAY08660
ID AAY08660 standard; protein; 383 AA.
XX
XX AAY08660;
XX
XX 09-AUG-1999 (first entry)
XX
XX W09927094 Seq ID 12.
XX
XX Transmembrane domain; human; nutrition; cytokine; cell differentiation;
KW immune stimulation; immune suppression; haematopoiesis; activin;

KW regulatory tissue growth; inhibin; chemostatic; chemokinetic;
 KW haemostatic; thrombolytic; tumour inhibitor; anti-inflammatory;
 XX gene therapy; screening.

OS Homo sapiens.

XX WO9927094-A2.

XX 03-JUN-1999.

XX 20-NOV-1998; 98WO-JP005238.

XX 25-NOV-1997; 97JP-00323129.

XX (SAGA) SAGAMI CHEM RES CENT.

XX (PROT-) PROTEGENE INC.

XX Kato S, Kimura T, Sekine S;

XX WPI; 1999-357835/30.

XX Novel proteins containing transmembrane domains, useful as anti-inflammatory, immune stimulators/suppressors and tissue growth compounds.

XX Disclosure; Page 87-89; 89pp; English.

XX This invention describes novel human transmembrane containing proteins and their encoding nucleic acids. Although no specific use is given for the proteins, they may have a range of activities selected from nutritional uses, cytokine and cell differentiation, immune stimulation/suppression, haematopoiesis regulatory, tissue growth, activin/inhibin, chemostatic/chemokinetic, haemostatic/thrombolytic, receptor/ligand, tumour inhibitor, anti-inflammatory and other undefined activities. The cDNAs can be utilized as probes for gene diagnosis and as gene sources for gene therapy. The cDNAs can also be used for large scale expression of proteins. The transformed cells can be used for detection of the corresponding ligands and for screening of novel low-molecular pharmaceuticals

XX Sequence 383 AA;

Query Match 97.6%; Score 2080; DB 2; Length 383;
 Best Local Similarity 100.0%; Pred. No. 2.8e-147; Indels 0; Gaps 0;
 Matches 383; Conservative 0; Mismatches 0;

QY	1	MAGIPGLLFLPFLCAVGVSPYAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
DB	1	MAGIPGLLFLPFLCAVGVSPYAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
QY	61	VSSSCGPOCHKTPLTYEAKQYLSYETLYANGSRTEQVGIYILSSGSGAQHRDGS	120
DB	61	VSSSCGPOCHKTPLTYEAKQYLSYETLYANGSRTEQVGIYILSSGSGAQHRDGS	120
QY	121	SGSKRRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTGLVAEKHVLTAACHIDG	180
DB	121	SGSKRRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTGLVAEKHVLTAACHIDG	180
QY	181	KTYVKGTKLVLGFLKPKFGDGGRGANDSTSAPEQMKFQWIRKVRTHVPKGIKNAND	240
DB	181	KTYVKGTKLVLGFLKPKFGDGGRGANDSTSAPEQMKFQWIRKVRTHVPKGIKNAND	240
QY	241	IGMDYDYLLELKKPKRPFMTKIGVSPAPKQLPGGRIFHSGYNDPGLNLYRFDCKDE	300
DB	241	IGMDYDYLLELKKPKRPFMTKIGVSPAPKQLPGGRIFHSGYNDPGLNLYRFDCKDE	300
QY	301	TYDLLYQCDAPGASGSGVYVEMWKRQOKWERKLIIGIFSGHWDNMGSPQDFNVAVR	360
DB	301	TYDLLYQCDAPGASGSGVYVEMWKRQOKWERKLIIGIFSGHWDNMGSPQDFNVAVR	360
QY	361	ITPLKTAQICYWIKGNLYDCREG	383
DB	361	ITPLKTAQICYWIKGNLYDCREG	383

RESULT 4

AY08657

XX ID AAY08657 standard; protein; 383 AA.

XX AC AAY08657;

XX DT 09-AUG-1999 (first entry)

XX DE Human transmembrane domain containing protein from clone HP10493.

XX Transmembrane domain; human; nutrition; cytokine; cell differentiation;
 KW immune stimulation; immune suppression; haematopoiesis; activin;
 KW regulatory tissue growth; inhibin; chemostatic; chemokinetic;
 KW haemostatic; thrombolytic; tumour inhibitor; anti-inflammatory;
 XX gene therapy; screening.

XX OS Homo sapiens.

XX WO9927094-A2.

XX 03-JUN-1999.

XX 20-NOV-1998; 98WO-JP005238.

XX 25-NOV-1997; 97JP-00323129.

XX (SAGA) SAGAMI CHEM RES CENT.

XX (PROT-) PROTEGENE INC.

XX Kato S, Kimura T, Sekine S;

XX WPI; 1999-357835/30.

XX N-PSDB; AAX77690, AAX77691.

XX Novel proteins containing transmembrane domains, useful as anti-inflammatory, immune stimulators/suppressors and tissue growth compounds.

XX Claim 1; Page 68-69; 89pp; English.

XX This invention describes novel human transmembrane containing proteins and their encoding nucleic acids. Although no specific use is given for the proteins, they may have a range of activities selected from nutritional uses, cytokine and cell differentiation, immune stimulation/suppression, haematopoiesis regulatory, tissue growth, activin/inhibin, chemostatic/chemokinetic, haemostatic/thrombolytic, receptor/ligand, tumour inhibitor, anti-inflammatory and other undefined activities. The cDNAs can be utilized as probes for gene diagnosis and as gene sources for gene therapy. The cDNAs can also be used for large scale expression of proteins. The transformed cells can be used for detection of the corresponding ligands and for screening of novel low-molecular pharmaceuticals

XX Sequence 383 AA;

Query Match 97.6%; Score 2080; DB 2; Length 383;
 Best Local Similarity 100.0%; Pred. No. 2.8e-147; Indels 0; Gaps 0;
 Matches 383; Conservative 0; Mismatches 0;

QY	1	MAGIPGLLFLPFLCAVGVSPYAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
DB	1	MAGIPGLLFLPFLCAVGVSPYAPWKPPTWPAAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
QY	61	VSSSCGPOCHKTPLTYEAKQYLSYETLYANGSRTEQVGIYILSSGSGAQHRDGS	120
DB	61	VSSSCGPOCHKTPLTYEAKQYLSYETLYANGSRTEQVGIYILSSGSGAQHRDGS	120
QY	121	SGSKRRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTGLVAEKHVLTAACHIDG	180
DB	121	SGSKRRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTGLVAEKHVLTAACHIDG	180

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QY 181 KTYVKGTKLRVGLFKPKFKDGGGANDSTSAMPEQMKFQWIRVKRTHVTPKGIKGNAND 240
Db 181 KTYVKGTKLRVGLFKPKFKDGGGANDSTSAMPEQMKFQWIRVKRTHVTPKGIKGNAND 240
QY 241 IGMDDYALLELKKPKRKFMKIGVSPPAKQLPGGRHFSGYDNDPRGNLVYRFFCDVKDE 300
Db 241 IGMDDYALLELKKPKRKFMKIGVSPPAKQLPGGRHFSGYDNDPRGNLVYRFFCDVKDE 300
QY 301 TYDLLYQQCDAQPGASGGYVYRMKRRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAQPGASGGYVYRMKRRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPKVAQICYWKIGNYLDREG 383
Db 361 ITPKVAQICYWKIGNYLDREG 383

RESULT 5
AAV13390
ID AAV13390 standard; protein; 383 AA.
AC AAV13390;
XX
XX
DT 25-JUN-1999 (first entry)
XX
DE Amino acid sequence of protein PRO307.
XX
KW Secreted protein; transmembrane protein; human; enterocolitis;
KW Zollinger-Ellison syndrome; gastrointestinal ulceration;
KW congenital microvillus atrophy; skin disease; cell growth;
KW abnormal keratinocyte differentiation; psoriasis; epithelial cancer;
KW Parkinson's disease; Alzheimer's disease; ALS; neuropathy; fibromodulin;
KW dermal scarring; Usher Syndrome; Atrophla areata; anti-thrombotic;
KW wound healing; tissue repair.
XX
XX Homo sapiens.
OS
XX
XX WO9914328-A2.
XX
XX 25-MAR-1999.
XX
XX 16-SEP-1998; 98WO-US019330.
XX
XX 17-SEP-1997; 97US-0059113P.
XX 17-SEP-1997; 97US-0059115P.
XX 17-SEP-1997; 97US-0059117P.
XX 17-SEP-1997; 97US-0059119P.
XX 17-SEP-1997; 97US-0059121P.
XX 17-SEP-1997; 97US-0059122P.
XX 17-SEP-1997; 97US-0059184P.
XX 18-SEP-1997; 97US-0059263P.
XX 18-SEP-1997; 97US-0059266P.
XX 15-OCT-1997; 97US-0062125P.
XX 17-OCT-1997; 97US-0062285P.
XX 17-OCT-1997; 97US-0062287P.
XX 21-OCT-1997; 97US-0063486P.
XX 24-OCT-1997; 97US-0062814P.
XX 24-OCT-1997; 97US-0062816P.
XX 24-OCT-1997; 97US-0063045P.
XX 24-OCT-1997; 97US-0063120P.
XX 24-OCT-1997; 97US-0063121P.
XX 24-OCT-1997; 97US-0063127P.
XX 24-OCT-1997; 97US-0063128P.
XX 27-OCT-1997; 97US-0063327P.
XX 27-OCT-1997; 97US-0063329P.
XX 28-OCT-1997; 97US-0063541P.
XX 28-OCT-1997; 97US-0063542P.
XX 28-OCT-1997; 97US-0063544P.
XX 28-OCT-1997; 97US-0063549P.
XX 28-OCT-1997; 97US-0063550P.
XX 28-OCT-1997; 97US-0063564P.
XX 29-OCT-1997; 97US-0063435P.
XX 29-OCT-1997; 97US-0063704P.

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PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
XX
XX (GETH ) GENENTECH INC.
XX
XX Wood WI, Gurney AL, Goddard A, Pennica D, Chen J, Yuan J;
PI
XX
XX WPI; 1999-229533/19.
DR N-PSDB; AAX52261.
XX
XX New isolated human genes and polypeptides used in, e.g. treatment of
PT gastrointestinal ulceration.
XX
XX Claim 12; Fig 96; 320pp; English.
XX
XX AAV13344-403 represent secreted and transmembrane human proteins. The
CC cDNA sequences are obtained from cDNA libraries, prepared from fetal
CC lung, fetal kidney, fetal brain, fetal liver and fetal retina. The
CC encoded polypeptides have specific uses based on their homology to known
CC polypeptides, e.g. PRO211 and PRO217 can be used for disorders associated
CC with the preservation and maintenance of gastrointestinal mucosa and the
CC repair of acute and chronic mucosal lesions (e.g. enterocolitis.
CC Zollinger-Ellison syndrome, gastrointestinal ulceration and congenital
CC microvillus atrophy), skin diseases associated with abnormal keratinocyte
CC differentiation (e.g. psoriasis, epithelial cancers such as lung squamous
CC cell carcinoma of the vulva and gliomas), potent effects on cell growth
CC and development, diseases related to growth or survival of nerve cells
CC including Parkinson's disease, Alzheimer's disease, ALS, neuropathies or
CC cancer. PRO265 can be used as for fibromodulin, e.g. for reducing dermal
CC scarring. PRO264 can be used as a target for anti-tumor drugs. PRO533 may
CC be used in the treatment of Usher Syndrome or Atrophla areata; PRO269 can
CC be used as an anti-thrombotic agent; PRO287 polypeptides and portions may
CC have therapeutic applications in wound healing and tissue repair; PRO317
CC can be used for treating problems of the kidney, uterus, endometrium,
CC blood vessels, or related tissue, e.g. in the heart of genital tract
XX
XX Sequence 383 AA;
SQ
Query Match 97.6%; Score 2080; DB 2; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.8e-147;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MAGIPGLLFLFFLLCAVQGVSPYSAPMKPTWPAYRLPVLPQSTLNLAKPFGAEAKLE 60
Db 1 MAGIPGLLFLFFLLCAVQGVSPYSAPMKPTWPAYRLPVLPQSTLNLAKPFGAEAKLE 60
QY 61 VSSSCGPOCHKGTPLPTYEAKQVLSYETLYANGSTETQVGIYILSSGDAQHRDSCS 120
Db 61 VSSSCGPOCHKGTPLPTYEAKQVLSYETLYANGSTETQVGIYILSSGDAQHRDSCS 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
QY 181 KTYVKGTKLRVGLFKPKFKDGGGANDSTSAMPEQMKFQWIRVKRTHVTPKGIKGNAND 240

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Db 181 KTVVGTQKLRVGLFKPKFGKGGRGANDSTSAPEQMKFQWIRKRVKTHVPKGIKGNAND 240
QY 241 IGMVDYDYLLELKKPKRKFPMKIGVSPPAKQLPGGRHFSGYDNDPRGNLVYRFDVKDE 300
Db 241 IGMVDYDYLLELKKPKRKFPMKIGVSPPAKQLPGGRHFSGYDNDPRGNLVYRFDVKDE 300
QY 301 TYDLLYQCCDAQPGASGSGVYVVMWKRQKQKWERKIIGIFSGHGWYDMNGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAQPGASGSGVYVVMWKRQKQKWERKIIGIFSGHGWYDMNGSPQDFNVAVR 360
QY 361 ITPKYAQICYWIKNYLDCCREG 383
Db 361 ITPKYAQICYWIKNYLDCCREG 383

RESULT 6
AAY88277
ID AAY88277 standard; protein; 383 AA.

AC AAY88277;
XX
DT 16-OCT-2000 (first entry)
XX
DE Human TANGO 186 protein.
XX
KW TANGO 180; TANGO 181; TANGO 182; TANGO 183; TANGO 184; TANGO 185;
KW TANGO 186; TANGO 188; TANGO 189; TANGO 215; TANGO 187; human; murine;
KW secreted protein; transmembrane protein; gene therapy; vaccine;
KW diagnosis; treatment; detection.
XX
OS Homo sapiens.
XX
PN WO200018904-A2.
XX
PD 06-APR-2000.
XX
PF 30-SEP-1999; 99WO-US022817.
XX
PR 30-SEP-1998; 98US-00164220.
PR 02-OCT-1998; 98US-00164169.
XX
PA (WILL-) MILLENNIUM BIOTHERAPEUTICS INC.
XX
PI Barnes TW;
XX
DR WPI; 2000-293144/25.
DR N-PSDB; AAA39945, AAA39946.

PT Isolated nucleic acids encoding TANGO polypeptides useful for preventing,
PT diagnosing and treating diseases associated with inappropriate protein
PT expression.
XX
XX Claim 9; Fig 13; 249pp; English.
PS
XX This invention describes novel human and murine nucleic acids encoding
CC TANGO polypeptides (which are either wholly secreted or transmembrane
CC proteins) which can be used for gene therapy and/or vaccination. The
CC peptides are designated TANGO 180 to TANGO 189 and TANGO 215. The nucleic
CC acids may be used to produce TANGO 180 to TANGO 189 and TANGO 215
CC polypeptides according to standard recombinant DNA methodologies. They
CC may also be used to detect and quantify the presence of TANGO nucleic
CC acids in a sample and therefore identify or diagnose diseases associated
CC with inappropriate TANGO expression (e.g. diseases related to over or
CC under expression of the polypeptides or the expression of inactive
CC polypeptides). The nucleic acids and the polypeptides they encode may be
CC used according to standard gene therapy protocols, to treat diseases
CC associated with inappropriate TANGO expression by supplementing a
CC patients own production of the polypeptide of to rectify mutations that
CC may result in expression of an abnormally active polypeptide. The
CC polypeptides may also be used to identify and produce agonists and
CC antagonists of TANGO expression and activity which may be used to
CC modulate TANGO related processes and diseases. The polypeptides are

CC particularly useful for use as antigens for producing antibodies to TANGO
CC proteins which may be used for inhibiting the activity of TANGO proteins.
CC They may also be used to detect and quantify the presence of TANGO
CC proteins in samples and therefore identify patients in whom the protein
CC is over- or under-expressed. This sequence represents the human TANGO 186
CC protein described in the method of the invention
XX

SQ Sequence 383 AA;

Query Match 97.6%; Score 2080; DB 3; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.8e-147;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYAPWKPWPAYRLPVVLPQSTLNLAKPDPFGAAKLE 60
Db 1 MAGIPGLLFLFLCAVGVSPYAPWKPWPAYRLPVVLPQSTLNLAKPDPFGAAKLE 60
QY 61 VSSSCGPQCHKTPLPTTVEEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDSGS 120
Db 61 VSSSCGPQCHKTPLPTTVEEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDSGS 120
QY 121 SGKSRKROIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEKHVLTAACIHG 180
Db 121 SGKSRKROIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEKHVLTAACIHG 180
QY 181 KTYVKGTKLRVGLFKPKFGKGGRGANDSTSAPEQMKFQWIRKRVKTHVPKGIKGNAND 240
Db 181 KTYVKGTKLRVGLFKPKFGKGGRGANDSTSAPEQMKFQWIRKRVKTHVPKGIKGNAND 240
QY 241 IGMVDYDYLLELKKPKRKFPMKIGVSPPAKQLPGGRHFSGYDNDPRGNLVYRFDVKDE 300
Db 241 IGMVDYDYLLELKKPKRKFPMKIGVSPPAKQLPGGRHFSGYDNDPRGNLVYRFDVKDE 300
QY 301 TYDLLYQCCDAQPGASGSGVYVVMWKRQKQKWERKIIGIFSGHGWYDMNGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAQPGASGSGVYVVMWKRQKQKWERKIIGIFSGHGWYDMNGSPQDFNVAVR 360
QY 361 ITPKYAQICYWIKNYLDCCREG 383
Db 361 ITPKYAQICYWIKNYLDCCREG 383

RESULT 7
AAY87270
ID AAY87270 standard; protein; 383 AA.

AC AAY87270;
XX
DT 11-MAY-2000 (first entry)
XX
DE Human signal peptide containing protein HSPP-47 SEQ ID NO:47.
XX
KW Human; signal peptide-containing protein; HSPP; diagnosis; cancer;
KW inflammation; cardiovascular disease; anticancer; anti-inflammatory;
KW antimicrobial; nootropic; neuroprotective; cardiovascular; hepatotropic;
KW antiasthmatic; gene therapy; cell proliferation; neurological disorder;
KW reproductive disorder; developmental disorder; arteriosclerosis;
KW cirrhosis; psoriasis; acquired immune deficiency syndrome; anaemia;
KW asthma; Crohn's disease; infection; Alzheimer's disease; schizophrenia;
KW Parkinson's disease; Huntington's disease; ovulatory defect;
KW muscular dystrophy.
XX
OS Homo sapiens.
XX
PN WO200000610-A2.
XX
PD 06-JAN-2000.
XX
PF 25-JUN-1999; 99WO-US014484.
XX
PR 26-JUN-1998; 98US-0090762P.
PR 31-JUL-1998; 98US-0094983P.
PR 01-OCT-1998; 98US-0102686P.

PR 11-DEC-1998; 98US-0112129P.
XX (INCY-) INCYTE PHARM INC.
XX Lal P, Tang YT, Gorgone GA, Corley NC, Guegler KJ, Baughn ME;
PI Akerblom IE, Au-Young J, Yue H, Patterson C, Reddy R, Hillman JL;
PI Bandman O;
XX WPI; 2000-160673/14.
DR N-PSDB; AA298155.
XX New human signal peptide-containing proteins useful in treatment,
PT prevention and diagnosis of e.g. cancer, inflammation and cardiovascular
PT disease.
XX Claim 1; Page 192-193; 327pp; English.
XX AA298109 to AA298242 encode AA298224 to AA298237 which represent the
CC human signal peptide-containing proteins HSP-1 to HSP-134. HSPs have
CC anticancer, anti-inflammatory, antimicrobial, nontoxic, hepatotropic,
CC neuroprotective, cardiovascular and antiasthmatic activities, and can be
CC used in gene therapy. HSPs can be used to treat or prevent disorders
CC associated with decreased activity or function of HSP. Antagonists of
CC HSP are used to treat or prevent disorders associated with increased
CC activity or function of HSP. Such diseases include cell proliferation
CC (including cancer), inflammation, cardiovascular, neurological,
CC reproductive or developmental disorders, (e.g. arteriosclerosis,
CC asthma, Crohn's disease, microbial or other infections, congestive or
CC ischaemic heart disease, Alzheimer's, Parkinson's or Huntington's
CC diseases, schizophrenia, ovulatory defects, muscular dystrophy). HSP
CC nucleic acids can be used for the recombinant production of HSP, for
CC detecting HSP in standard hybridization and amplification assays (for
CC diagnosis and monitoring), in gene therapy, as antisense, triplex-forming
CC or ribozyme therapeutics, for detecting related sequences or genetic
CC variations, and for chromosomal mapping. HSP are also used to raise
CC specific antibodies (Ab) and to screen for agonists and antagonists
CC (potential therapeutic agents). Ab are used to diagnose, or monitor, HSP
CC -related diseases (in usual immunoassays), as therapeutic antagonists, in
CC competitive drug screens, and for purification of HSP from natural
CC sources
XX SQ Sequence 383 AA;
Query Match 97.6%; Score 2080; DB 3; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.8e-147;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MAGIPGLLFLPFLCAVGVSPYSAPWKTWPAYRLPVVLQSTLNLAKPFGAEAKLE 60
DB 1 MAGIPGLLFLPFLCAVGVSPYSAPWKTWPAYRLPVVLQSTLNLAKPFGAEAKLE 60
QY 61 VSSSCGPOCHKPTLPTYEAKQYLSYETLYANGSTETQVGIYLLSSGDAQHRDSSG 120
DB 61 VSSSCGPOCHKPTLPTYEAKQYLSYETLYANGSTETQVGIYLLSSGDAQHRDSSG 120
QY 121 SGKSRKRQIYGVDSRFSIFGKDFLNLNYPFSTSVKLSLSTGCTGLVAEKHLVLAACHIDG 180
DB 121 SGKSRKRQIYGVDSRFSIFGKDFLNLNYPFSTSVKLSLSTGCTGLVAEKHLVLAACHIDG 180
QY 181 KTVVGTQKLRVFLKPKFKDGGGRANDSTSAMPEQMKFQWIRVVRKTHVPKGMIGNAND 240
DB 181 KTVVGTQKLRVFLKPKFKDGGGRANDSTSAMPEQMKFQWIRVVRKTHVPKGMIGNAND 240
QY 241 IGMVDYALLELKKPKRPMKIGVSPRAKQIPGRIHFSGVNDPBGNLVRFCDVKDE 300
DB 241 IGMVDYALLELKKPKRPMKIGVSPRAKQIPGRIHFSGVNDPBGNLVRFCDVKDE 300
QY 301 TYDLLYQQCDAQPGASGSGVYVVMWKRQCKQKWERKLIIGFSGHQWDMNGSPQDFNVAVR 360
DB 301 TYDLLYQQCDAQPGASGSGVYVVMWKRQCKQKWERKLIIGFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPLKYAQICYWIKGNLYDCREG 383

DB 361 ITPLKYAQICYWIKGNLYDCREG 383
RESULT 8
AA298227
ID AA298227 standard; protein; 383 AA.
XX AA298227;
XX 22-FEB-2000 (first entry)
DT DE A bone marrow secreted protein designated BMS192.
XX Bone marrow secreted protein; bone marrow stromal cell; cytokine;
KW cell proliferation; cell differentiation; hematopoiesis; anaemia;
KW myeloid cell deficiency; lymphoid cell deficiency; myeloid cell;
KW erythroid progenitor cell; colony stimulating factor; granulocyte;
KW monocyte; macrophage; myelo-suppression; megakaryocyte; platelet;
KW platelet disorder; thrombocytopenia; hematopoietic stem cell;
KW stem cell disorder; aplastic anaemia; bone differentiation;
KW paroxysmal nocturnal hemoglobinuria; bone growth; cartilage; tendon;
KW ligament; nerve; wound healing; tissue repair; burn; incision; ulcer;
KW bone fracture; cartilage damage; artificial joint.
XX Homo sapiens.
OS FH Key Location/Qualifiers
FT Peptide 1..19 /note= "signal peptide"
FT W099333979-A2.
XX 08-JUL-1999.
XX 18-DEC-1999; 98WO-US027008.
XX 30-DEC-1997; 97US-0068958P.
PR 24-SEP-1998; 98US-0101603P.
PR 30-SEP-1998; 98US-0102540P.
XX (CHIR) CHIRON CORP.
XX Lin H, Cao L;
XX WPI: 2000-038344/03.
DR N-PSDB; AA236233.
XX New isolated human polynucleotide and secreted proteins can induce
PT production of other cytokines in certain cell populations.
XX Claim 2; Page 81-82; 120pp; English.
XX AA29822-43 represent bone marrow secreted proteins of human bone marrow
CC stromal cells. The proteins can exhibit cytokine, cell proliferation, or
CC cell differentiation activity (either inducing or inhibiting). They can
CC be used to support colony forming cells or factor-dependent cell lines,
CC to regulate hematopoiesis, and to treat myeloid or lymphoid cell
CC deficiencies. In addition, they may be used to support the growth and
CC proliferation of erythroid progenitor cells, and to treat various
CC anaemias. They can have colony stimulating factor (CSF) activity and can
CC be used to support the growth and proliferation of myeloid cells such as
CC granulocytes, monocytes or macrophages, to prevent or treat myelo-
CC suppression, to support the growth and proliferation of megakaryocytes
CC and platelets, thereby allowing prevention or treatment of platelet
CC disorders such as thrombocytopenia, to support the growth and
CC proliferation of hematopoietic stem cells, either in place of or in
CC conjunction with platelet transfusions, to treat stem cell disorders,
CC such as aplastic anaemia and paroxysmal nocturnal hemoglobinuria, or to
CC repopulate the stem cell compartment after irradiation or chemotherapy.
CC They can be used for growth or differentiation of bone, cartilage, and
CC tendon, ligament, or nerve tissue, as well as for wound healing and
CC tissue repair and replacement, and in the treatment of burns, incisions

CC and ulcers, to induce cartilage and/or bone growth in circumstances where
CC bone is not normally formed and thus have an application in healing bone
CC fractures and cartilage damage or defects, prophylactic use in fracture
CC reduction and also in the improved fixation of artificial joints
XX
SQ Sequence 383 AA;

Query Match 97.6%; Score 2080; DB 3; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.8e-147;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPOCHKTPLTYEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDGS 120
DB 61 VSSSCGPOCHKTPLTYEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180

QY 181 KTYVKGTKRLRGFLKPKFKDGGGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240
DB 181 KTYVKGTKRLRGFLKPKFKDGGGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240

QY 241 IGMVDYDYLLELKKPKRKFMIKIGVSPAPKQLPGGRHPSGYNDNDRPGLNVRFCVDKDE 300
DB 241 IGMVDYDYLLELKKPKRKFMIKIGVSPAPKQLPGGRHPSGYNDNDRPGLNVRFCVDKDE 300

QY 301 TYDILLYQCCDAQPGASGSGVYVVMKROQKWKIIGIFSGHWDVMDNGSPQDFNVAVR 360
DB 301 TYDILLYQCCDAQPGASGSGVYVVMKROQKWKIIGIFSGHWDVMDNGSPQDFNVAVR 360

QY 361 ITPLKYAQICYWIKGNLYDCREG 383
DB 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 9
AAB25618
ID AAB25618 standard; protein; 383 AA.

XX AAB25618;
XX
XX
XX 21-NOV-2000 (first entry)
XX
XX Protein encoded by human secreted protein gene #10.

DE Secreted protein; immunosuppressant; anti-inflammatory; antiarthritic;
KW antirheumatic; dermatological; antiproliferative; antiarteriosclerotic;
KW anticancer; vulnary; antiviral; antibacterial; antifungal;
KW immune disorder; Addison's disease; hyperproliferative disorders e.g. coronary
KW multiple sclerosis; inflammatory disorder; rheumatoid arthritis; dermatitis;
KW Crohn's disease; nephritis; hyperproliferative disorder;
KW cardiovascular disorder; coronary arteriosclerosis; myocarditis; cancer;
KW melanoma; lymphoma; wound healing; human; chromosome 12.

XX Homo sapiens.
XX
XX WO200029435-A1.
XX
XX 25-MAY-2000.
XX
XX 27-OCT-1999; 99WO-US025031.
XX
XX 28-OCT-1998; 98US-0105971P.
XX
XX (HUMA-) HUMAN GENOME SCI INC.
XX
XX Ni J, Ruben SM, Olesen HS, Young PE, Kenny JJ, Moore PA, Wei Y;
XX Greene JM;

XX WPI; 2000-387742/33;
XX Isolated nucleic acid molecules encoding human secreted proteins are used
PT for the prevention, amelioration and treatment of autoimmune,
PT inflammatory, hyperproliferative and cardiovascular disorders, cancer,
PT wounds, and infectious diseases.
XX
XX Disclosure; Page 169; 803pp; English.
XX

CC The present invention relates to 12 secreted human proteins and the
CC nucleotide sequences encoding them. The polynucleotide sequences given in
CC AAB0606-A80623 encode the 12 secreted protein sequences given in
CC AAB25576-B25593. The human secreted proteins have various activities
CC dependent on the tissues in which they are expressed. Examples of the
CC activities of the proteins include: immunosuppressant; anti-inflammatory;
CC antirheumatic; antirheumatic; dermatological; antiproliferative;
CC antiarteriosclerotic; anticancer; vulnary; antiviral; antibacterial;
CC and antifungal activity. The proteins, polypeptides, agonists and
CC antagonists may be used to treat prevent and/or diagnose various disease,
CC disorders and conditions examples of which include: immune disorders e.g.
CC Addison's disease, rheumatoid arthritis, dermatitis, and multiple
CC sclerosis; inflammatory disorders e.g. inflammatory bowel disease,
CC Crohn's disease and nephritis; hyperproliferative disorders such as
CC parotiteinaemia and purpura; cardiovascular disorders e.g. coronary
CC arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma.
CC proteins and polynucleotide sequences may also be used in wound healing
CC and the treatment of infectious diseases. The human secreted protein gene
CC #10 and protein sequences are represented in sequences AAB0615 and
CC AAB25585. Secreted protein gene #10 is located on chromosome 12.
CC Sequences AAB25616-B25618 represent alternative secreted protein #10
CC sequences and AAB06629-A80676 represent genes which are related to the
CC secreted protein gene#10
XX Sequence 383 AA;

Query Match 97.6%; Score 2080; DB 3; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.8e-147;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPOCHKTPLTYEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDGS 120
DB 61 VSSSCGPOCHKTPLTYEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180

QY 181 KTYVKGTKRLRGFLKPKFKDGGGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240
DB 181 KTYVKGTKRLRGFLKPKFKDGGGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240

QY 241 IGMVDYDYLLELKKPKRKFMIKIGVSPAPKQLPGGRHPSGYNDNDRPGLNVRFCVDKDE 300
DB 241 IGMVDYDYLLELKKPKRKFMIKIGVSPAPKQLPGGRHPSGYNDNDRPGLNVRFCVDKDE 300

QY 301 TYDILLYQCCDAQPGASGSGVYVVMKROQKWKIIGIFSGHWDVMDNGSPQDFNVAVR 360
DB 301 TYDILLYQCCDAQPGASGSGVYVVMKROQKWKIIGIFSGHWDVMDNGSPQDFNVAVR 360

QY 361 ITPLKYAQICYWIKGNLYDCREG 383
DB 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 10
AAB25592
ID AAB25592 standard; protein; 383 AA.
XX

AA25592;
21-NOV-2000 (first entry)
Protein encoded by human secreted protein gene #10 clone HUSQ05.
Secreted protein; immunosuppressant; anti-inflammatory; antiarthritic;
antirheumatic; dermatological; antiproliferative; antiarthritic;
antitumor; vulvar; antiviral; antibacterial; antifungal;
immune disorder; Addison's disease; rheumatoid arthritis; dermatitis;
multiple sclerosis; inflammatory disorder; inflammatory bowel disease;
Crohn's disease; nephritis; hyperproliferative disorder;
cardiovascular disorder; coronary arteriosclerosis; myocarditis; cancer;
melanoma; lymphoma; wound healing; human; chromosome 12.
Homo sapiens.
WO200029435-A1.
25-MAY-2000.
27-OCT-1999; 99WO-US025031.
28-OCT-1998; 98US-0105971P.
(HUMA-) HUMAN GENOME SCI INC.
Ni J, Ruben SM, Olsen HS, Young PE, Kenny JJ, Moore PA, Wei Y;
Greene JM;
WPI; 2000-387742/33.
Isolated nucleic acid molecules encoding human secreted proteins are used
for the prevention, amelioration and treatment of autoimmune,
inflammatory, hyperproliferative and cardiovascular disorders, cancer,
wounds, and infectious diseases.
Claim 1; Page 684-685; 803pp; English.
The present invention relates to 12 secreted human proteins and the
nucleotide sequences encoding them. The polynucleotide sequences given in
AA80606-A80623 encode the 12 secreted protein sequences given in
AA25576-B25593. The human secreted proteins have various activities
dependent on the tissues in which they are expressed. Examples of the
activities of the proteins include: immunosuppressant; anti-inflammatory;
antiarthritic; antirheumatic; dermatological; antiproliferative;
antiarteriosclerotic; anticancer; vulvar; antiviral; antibacterial;
and antifungal activity. The proteins, polypeptides, agonists and
antagonists may be used to treat prevent and/or diagnose various disease,
disorders and conditions examples of which include: immune disorders e.g.
Addison's disease, rheumatoid arthritis, dermatitis, and multiple
sclerosis; inflammatory disorders e.g. inflammatory bowel disease,
Crohn's disease and nephritis; hyperproliferative disorders such as
paraproteinemia and purpura; cardiovascular disorders e.g. coronary
arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The
proteins and polynucleotide sequences may also be used in wound healing
and the treatment of infectious diseases. The human secreted protein gene
#10 and protein sequences are represented in sequences AA80615 and
AA80585. Secreted protein gene #10 is located on chromosome 12.
Sequences AA805618-B25618 represent alternative secreted protein #10
sequences and AA80669-A80676 represent alternative genes which are related to the
secreted protein gene#10
Sequence 383 AA;
Query Match 97.6%; Score 2080; DB 3; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.8e-147;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
1 MAGIPGLLLFLLCAVGVSPYAPWPTWPAVRLPVVLPQSTNLAKPFGAEAKLE 60
1 MAGIPGLLLFLLCAVGVSPYAPWPTWPAVRLPVVLPQSTNLAKPFGAEAKLE 60

QY 61 VSSCGPQCHKGTPLPTVEEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDGS 120
DB 61 VSSCGPQCHKGTPLPTVEEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDGS 120
QY 121 SKSRRKQIYGYDSRESIFGKDFLLNYPFSTSVKLSGCTGLVAEKHVLTAACHIDG 180
DB 121 SKSRRKQIYGYDSRESIFGKDFLLNYPFSTSVKLSGCTGLVAEKHVLTAACHIDG 180
QY 181 KTYVKGTKLVRGFLKPKFKDGGRGANDSTAMPQMKFQWIRVVRKTHVPKMGWKNAND 240
DB 181 KTYVKGTKLVRGFLKPKFKDGGRGANDSTAMPQMKFQWIRVVRKTHVPKMGWKNAND 240
QY 241 IGMDYDVALLELKPKHKKFKMIGVSPPAKQIPGGRIFHPSGYDNDPFGNLFVFCVKDE 300
DB 241 IGMDYDVALLELKPKHKKFKMIGVSPPAKQIPGGRIFHPSGYDNDPFGNLFVFCVKDE 300
QY 301 TYDLLYQCCDAQPGASGGVYVMWKRQOQKWERKIIGIFSGHQVDMNGSPQDFNVAVR 360
DB 301 TYDLLYQCCDAQPGASGGVYVMWKRQOQKWERKIIGIFSGHQVDMNGSPQDFNVAVR 360
QY 361 ITPLKYAQICYWIKGNLYLDCREG 383
DB 361 ITPLKYAQICYWIKGNLYLDCREG 383
RESULT 11
ADC78573
ID ADC78573 standard; protein; 383 AA.
XX
AC ADC78573;
XX
DT 01-JAN-2004 (first entry)
XX
DE Human PRO307 protein.
XX
KW antinflammatory; antiulcer; cytostatic; antipsoriatic; antiparkinsonian;
nootropic; neuroprotective; vasotropic; chemotactic; angiogenic;
neurotrophic; osteopathic; antiasthmatic; antiarthritic; antirheumatic;
antiarteriosclerotic; cardiac; antidiabetic; cerebroprotective;
thrombolytic; immunomodulator; enterocolitis; Zollinger-Ellison syndrome;
gastrointestinal ulceration; psoriasis; cancer; Parkinson's disease;
Alzheimer's; ALS; neuropathy; dermal scarring; wound healing;
nerve repair; thrombosis; bone; cartilage formation; angiogenesis;
asthma; rheumatoid arthritis; multiple sclerosis; inflammatory disorder;
atherosclerosis; cardiac injury; infertility; premature aging; AIDS;
diabetes; stroke; gene therapy; transgenic; PRO; human.
XX Homo sapiens.
OS
XX WO200015796-A2.
XX
XX PD 23-MAR-2000.
XX
XX PF 15-SEP-1999; 99WO-US021090.
XX
XX PR 16-SEP-1998; 98WO-US019330.
XX
XX PA (GETH) GENENTECH INC.
XX
XX PI Chen J, Goddard A, Gurney AL, Hillan K, Pennica D, Wood WI;
PI Yuan J;
XX
XX DR WPI; 2000-271434/23.
XX
XX DR N-PSDB; ADC78572.
XX
XX PT Novel nucleic acids encoding secreted and transmembrane polypeptides with
PT homology, e.g. to growth and cancer-associated antigens.
XX
XX PS Claim 12; SEQ ID NO 261; 355pp; English.
XX
XX CC The invention relates to a novel nucleic acid encoding a PRO polypeptide.
CC The polypeptides and polynucleotides of the invention may be useful as
CC research tools and as therapeutics for treating enterocolitis, Zollinger-

CC Ellison syndrome, gastrointestinal ulceration, psoriasis, cancer,
 CC Parkinson's disease, Alzheimer's disease, ALS, neuropathies, dermal
 CC scarring and wound healing, nerve repair, thrombosis, bone and/or
 CC cartilage formation, angiogenesis, asthma, rheumatoid arthritis, multiple
 CC sclerosis, inflammatory disorders, atherosclerosis, cardiac injury,
 CC infertility, premature aging, AIDS, diabetes complications and stroke.
 CC The molecules may also be utilised during gene therapy procedures and
 CC transgenic animal production. The current sequence is that of the human
 CC PRO protein of the invention.
 XX
 SQ Sequence 383 AA;
 Query Match 97.6%; Score 2080; DB 3; Length 383;
 Best Local Similarity 100.0%; Pred. No. 2.8e-147;
 Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
 DB 1 MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
 QY 61 VSSSCGPOCHKGTPPTYEAKQYLSYETLYANGSRTETQVGIYILSSGGAQHRS 120
 DB 61 VSSSCGPOCHKGTPPTYEAKQYLSYETLYANGSRTETQVGIYILSSGGAQHRS 120
 QY 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
 DB 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
 QY 181 KTYVKGTKLVRGFLKPKFKDGGGRANDSTSAMPEQMKFQIRVKTHTVPGWIKGNAND 240
 DB 181 KTYVKGTKLVRGFLKPKFKDGGGRANDSTSAMPEQMKFQIRVKTHTVPGWIKGNAND 240
 QY 241 IGMVDYALLELKKPHKPKMKIGVSPAPKQIPGRIHFSGYNDNRPNLVYRFDVKDE 300
 DB 241 IGMVDYALLELKKPHKPKMKIGVSPAPKQIPGRIHFSGYNDNRPNLVYRFDVKDE 300
 QY 301 TYDLYVQCCDAQPGASGVVYVMWKRQOQKWERKLIIGIFSGHVDMMGSPQDENVAVR 360
 DB 301 TYDLYVQCCDAQPGASGVVYVMWKRQOQKWERKLIIGIFSGHVDMMGSPQDENVAVR 360
 QY 361 ITPKVAQICWIKGNVLDREG 383
 DB 361 ITPKVAQICWIKGNVLDREG 383
 RESULT 12
 AAB80258
 ID AAB80258 standard; protein; 383 AA.
 XX
 AC AAB80258;
 XX
 DT 24-APR-2001 (first entry)
 DE Human PRO307 protein.
 XX
 KW Human; PRO; dermatological; antipsoriatic; cytostatic; antiinflammatory;
 KW antiparkinsonian neurotropic; neuroprotective; vulnerary; cardiac;
 KW antiangiogenic; vasotrophic; antiasthmatic; antirheumatic; cancer;
 KW antiarthritic; antifertility; antidiabetic; antiviral; diabetes;
 KW ophthalmological; gene therapy; skin disease; gastrointestinal disorder;
 KW ischaemia; inflammation.
 XX
 OS Homo sapiens.
 XX
 PN WO200104311-A1.
 XX
 PD 18-JAN-2001.
 XX
 PF 22-FEB-2000; 2000WO-US0004414.
 XX
 PR 07-JUL-1999; 99US-0143048P.
 PR 26-JUL-1999; 99US-0145698P.
 PR 28-JUL-1999; 99US-0146222P.

PR 08-SEP-1999; 99WO-US020594.
 PR 13-SEP-1999; 99WO-US020944.
 PR 15-SEP-1999; 99WO-US021090.
 PR 15-SEP-1999; 99WO-US021547.
 PR 05-OCT-1999; 99WO-US023089.
 PR 29-NOV-1999; 99WO-US028214.
 PR 30-NOV-1999; 99WO-US028313.
 PR 02-DEC-1999; 99WO-US028564.
 PR 02-DEC-1999; 99WO-US028565.
 PR 16-DEC-1999; 99WO-US030095.
 PR 20-DEC-1999; 99WO-US030911.
 PR 20-DEC-1999; 99WO-US030999.
 PR 05-JAN-2000; 2000WO-US000219.
 XX
 XX (GETH) GENENTECH INC.
 XX
 PI Ashkenazi AJ, Botstein D, Deenoyers L, Eaton DL, Ferrara N;
 PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
 PI Godowski PJ, Grimaldi CJ, Gurney AL, Hillan KJ, Kljavin IJ;
 PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
 PI Williams PM, Wood WI;
 XX
 DR WPI; 2001-081051/09.
 DR N-PSDB; AAF72419.
 XX
 XX Sixty one nucleic acids encoding PRO polypeptides which are useful in the
 XX treatment of skin diseases (e.g. psoriasis), cancers (e.g. lung squamous
 XX cell carcinoma) and neurodegenerative diseases (e.g. Alzheimer's
 XX disease).
 PS Claim 1; Fig 96; 393pp; English.
 XX
 XX The present sequence is one of sixty one novel secreted and transmembrane
 XX PRO polypeptides. The PRO polypeptides are useful for treating skin
 XX diseases (e.g. psoriasis), cancers (e.g. lung squamous cell carcinoma),
 XX gastrointestinal disorders (e.g. enterocolitis), neurodegenerative
 XX diseases (e.g. Alzheimer's disease, Parkinson's disease), wound repair,
 XX cardiovascular disorders (e.g. endometrial bleeding angiogenesis,
 XX ischaemia such as coronary ischaemia, atherosclerosis), inflammatory
 XX disorders (e.g. asthma, rheumatoid arthritis, multiple sclerosis),
 XX infertility, AIDS and diabetes and retinal disorders such as retinitis
 XX pigmentosa. The PRO nucleic acids have applications in molecular
 XX biology, including use as hybridization probes, and in chromosome and
 XX gene mapping
 XX
 SQ Sequence 383 AA;
 Query Match 97.6%; Score 2080; DB 4; Length 383;
 Best Local Similarity 100.0%; Pred. No. 2.8e-147;
 Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
 DB 1 MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
 QY 61 VSSSCGPOCHKGTPPTYEAKQYLSYETLYANGSRTETQVGIYILSSGGAQHRS 120
 DB 61 VSSSCGPOCHKGTPPTYEAKQYLSYETLYANGSRTETQVGIYILSSGGAQHRS 120
 QY 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
 DB 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
 QY 181 KTYVKGTKLVRGFLKPKFKDGGGRANDSTSAMPEQMKFQIRVKTHTVPGWIKGNAND 240
 DB 181 KTYVKGTKLVRGFLKPKFKDGGGRANDSTSAMPEQMKFQIRVKTHTVPGWIKGNAND 240
 QY 241 IGMVDYALLELKKPHKPKMKIGVSPAPKQIPGRIHFSGYNDNRPNLVYRFDVKDE 300
 DB 241 IGMVDYALLELKKPHKPKMKIGVSPAPKQIPGRIHFSGYNDNRPNLVYRFDVKDE 300
 QY 301 TYDLYVQCCDAQPGASGVVYVMWKRQOQKWERKLIIGIFSGHVDMMGSPQDENVAVR 360

Db 301 TYDLLYQQCDAQPGASSGVYVYRMWKRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
 QY 361 ITPLKYAQICYWIKGNLYDCREG 383
 Db 361 ITPLKYAQICYWIKGNLYDCREG 383
 RESULT 13
 AAB48974
 ID AAB48974 standard; protein; 383 AA.
 AC AAB48974;
 XX
 XX 27-MAR-2001 (first entry)
 XX
 DE Human Zsig13 variant #3, SEQ ID NO:18.
 XX
 KW Human Zsig13; serine protease; chromosome 11q22.1; elastase homologue;
 KW glutamyl endopeptidase homologue; factor X homologue; trypsin homologue;
 KW trypsinogen homologue; mast cell protease homologue;
 KW collagenase homologue; protein degradation; food processing; brewing;
 KW alcohol production; laundry detergent component.
 XX
 OS Homo sapiens.
 XX
 PN US6153420-A.
 XX
 PD 28-NOV-2000.
 XX
 PF 04-MAY-1998; 98US-00072384.
 XX
 PR 24-APR-1997; 97US-0044185P.
 PR 17-APR-1998; 98US-00062142.
 XX
 PA (ZYMO) ZYMOGENETICS INC.
 XX
 XX Sheppard PO;
 PI
 XX
 XX WPI; 2001-060090/07.
 DR N-PSDB; AAC91784.
 XX
 PT New isolated serine protease (designated Zsig13), useful in industrial
 PT processes to degrade unwanted proteins or alter the characteristics of
 PT protein-containing composition, as well as in industrial applications
 PT (e.g. brewing).
 XX
 XX Claim 1; Col 41-44; 26pp; English.
 XX
 CC The invention relates to human Zsig13 proteins (AAB48972-B48974), and to
 CC DNA encoding them (AAC91782-C91784). The invention also relates to
 CC expression vectors and host cells comprising a human Zsig13 DNA, and the
 CC recombinant production of a human Zsig13 protein or its precursor. Zsig13
 CC is a serine protease, and has significant homology to Bacillus
 CC licheniformis glutamyl endopeptidase, human clotting factor X, human
 CC elastase, rat mast cell protease, Streptomyces griseus trypsin, bovine
 CC trypsinogen, and Hypoderma lineatum collagenase. The gene encoding human
 CC Zsig13 is located on chromosome 11q22.1. Zsig13 is useful in industrial
 CC processes to degrade unwanted proteins or alter the characteristics of
 CC protein-containing compositions. It may also be used in industrial
 CC applications in which proteases are utilised, including food processing,
 CC brewing and alcohol production, and as a component of a laundry
 CC detergent. The present sequence represents a human Zsig13 variant
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 Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGTLVAEKHVLTAACHIDG 180
 QY 181 KTYVVGTKLRVGFLLKPKFKDGGRGANDSTSAAMPQMKFQWIRVKTHTVPKGIKGNAND 240
 Db 181 KTYVVGTKLRVGFLLKPKFKDGGRGANDSTSAAMPQMKFQWIRVKTHTVPKGIKGNAND 240
 QY 241 IGMWDYDIALLELKPKHKKFKMKIGVSPPAKQIPGGRIHPSGYDNDPGLNLYRFFCDVKDE 300
 Db 241 IGMWDYDIALLELKPKHKKFKMKIGVSPPAKQIPGGRIHPSGYDNDPGLNLYRFFCDVKDE 300
 QY 301 TYDLLYQQCDAQPGASSGVYVYRMWKRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
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 ID AAU29048 standard; protein; 383 AA.
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 AC AAU29048;
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 DT 18-DEC-2001 (first entry)
 XX
 DE Human PRO polypeptide sequence #25.
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 KW PRO polypeptide; mammal; tumour; cancer; human; cattle; horse; sheep;
 KW dog; cat; pig; goat; rabbit; tumour necrosis factor alpha; TNF-alpha;
 KW blood; chondrocyte cell; cell proliferation; cell differentiation; colon;
 KW adrenal; lung; breast; prostate; rectum; cervix; liver; genetic disorder.
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 OS Homo sapiens.
 XX
 PN WO200168848-A2.
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 PD 20-SEP-2001.
 XX
 PF 28-FEB-2001; 2001WO-US006520.
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 PR 01-MAR-2000; 2000WO-US005601.
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 PR 01-DEC-2000; 2000WO-US032678.
 PR 20-DEC-2000; 2000WO-US034956.
 XX
 PA (GETH) GENENTECH INC.
 XX
 XX Baker KP, Chen J, Desnoyers L, Goddard A, Godowski PJ, Gurney AL;
 PI Pan J, Smith V, Watanabe CK, Wood WI, Zhang Z;
 XX
 XX WPI; 2001-602746/68.
 DR N-PSDB; AAS45949.
 DR
 XX Novel nucleic acids encoding PRO polypeptides, used to diagnose the
 PT presence of tumors, such as prostate and breast tumors, in mammals and to
 PT screen for modulators of the compounds.
 XX
 XX Claim 11; Fig 50; 774pp; English.
 PS
 XX Sequences AAU29328-AAU29328 represent PRO polypeptides of the invention.
 CC The PRO polypeptides and their associated nucleic acids can be used to
 CC detect the presence of a tumour in a mammal by comparing the level of
 CC expression of a PRO polypeptide in a test sample of cells from the animal
 CC and a control sample of normal cells, whereby a higher level of
 CC expression in the test sample indicates the presence of a tumour in the
 CC mammal. Mammals include dogs, cats, cattle, horses, sheep, pigs, goats
 CC and rabbits but are preferably human. The polypeptides can be used to
 CC stimulate tumour necrosis factor (TNF) alpha release from human blood,
 CC when contacted with it. A specific polypeptide can be used to stimulate
 CC the proliferation or differentiation of chondrocyte cells. The PRO
 CC proteins can be used to determine the presence of tumours and also
 CC susceptibility to tumour development, particularly adrenal, lung, colon,
 CC breast, prostate, cervical or liver tumours, in mammalian
 CC subjects. The oligonucleotide probes specific for the PRO nucleic acids
 CC can be used for genetic analysis of individuals with genetic disorders
 XX
 SQ Sequence 383 AA;
 Query Match 97.6%; Score 2080; DB 4; Length 383;
 Best Local Similarity 100.0%; Pred. No. 2.8e-147;
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 DB 121 SGKSRKRIQYGVDSRFSFGKDFLNNYPTSVKLTGCTGLVAEKHVLTAACHIHG 180
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 AC ABUS8424;
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 DT 15-APR-2003 (first entry)
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 DE Human PRO polypeptide #25.
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 KW dog; cat; cow; horse; sheep; pig; goat; rabbit; ADAPT;
 KW antibody-dependent enzyme mediated prodrg therapy.
 XX
 OS Homo sapiens.
 XX
 PN US2003027272-A1.
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 PD 06-FEB-2003.
 XX
 PF 21-JUN-2002; 2002US-00176492.
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GenCore version 5.1.6
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OM protein - protein search, using sw model

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Listing first 45 summaries

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39	120	5.6	284	4	US-10-042-091A-7	Sequence 7, Appl
40	117	5.5	437	1	US-08-487-037-2	Sequence 2, Appl
41	115.5	5.4	241	3	US-08-944-483-59	Sequence 59, Appl
42	114	5.3	228	3	US-08-944-483-44	Sequence 44, Appl
43	114	5.3	253	6	5223425-8	Patent No. 5223425
44	114	5.3	253	6	5223425-8	Patent No. 5223425
45	113	5.3	238	6	5223425-5	Patent No. 5223425

ALIGNMENTS

RESULT 1
US-09-072-384-15
; Sequence 15, Application US/09072384
; Patent No. 6153420
; GENERAL INFORMATION:
; APPLICANT: Sheppard, Paul O.
; TITLE OF INVENTION: SERINE PROTEASE POLYPEPTIDES
; TITLE OF INVENTION: AND MATERIALS AND METHODS FOR MAKING THEM
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: ZymoGenetics, Inc.
; STREET: 1201 Eastlake Avenue East
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98102
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/072,384
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Parker, Gary E
; REGISTRATION NUMBER: 31,648
; REFERENCE/DOCKET NUMBER: 97-16CI
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-442-6673
; TELEFAX: 206-442-6678
; TELEX:
; INFORMATION FOR SEQ ID NO: 15:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 392 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FRAGMENT TYPE: internal
; FEATURE:
; NAME/KEY: Signal Sequence
; LOCATION: 1..19
; OTHER INFORMATION:
US-09-072-384-15

Query Match 100.0%; Score 2131; DB 3; Length 392;
Best Local Similarity 100.0%; Pred. No. 1.5e-225;
Matches 392; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPQCHKGTPPTVEEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRS DGS 120
DB 61 VSSSCGPQCHKGTPPTVEEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRS DGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEKHVLTAACHIDG 180
DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEKHVLTAACHIDG 180

QY 181 KTVVKGTKLRVGLFKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTHTVPKGIKNAND 240
DB 181 KTVVKGTKLRVGLFKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTHTVPKGIKNAND 240

QY 241 IGMDDYDYLLELKKPKFKMKGVSPPAKQLPGGRIHFSGYNDPRGNLVYRFCVDKDE 300
DB 241 IGMDDYDYLLELKKPKFKMKGVSPPAKQLPGGRIHFSGYNDPRGNLVYRFCVDKDE 300

QY 301 TYDLLYQCCDAQPGASGSGVYVVMKRWKQKWKIIGIFSGHQWDMNGSPQDFNVAVR 360
DB 301 TYDLLYQCCDAQPGASGSGVYVVMKRWKQKWKIIGIFSGHQWDMNGSPQDFNVAVR 360

QY 361 ITPLKYAQICYWKGNLYDCREGDTVFPPGNS 392
DB 361 ITPLKYAQICYWKGNLYDCREGDTVFPPGNS 392

RESULT 2

US-09-072-384-2
; Sequence 2, Application US/09072384
; Patent No. 6153420
; GENERAL INFORMATION:
; APPLICANT: Sheppard, Paul O.
; TITLE OF INVENTION: SERINE PROTEASE POLYPEPTIDES
; TITLE OF INVENTION: AND MATERIALS AND METHODS FOR MAKING THEM
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: ZymoGenetics, Inc.
; STREET: 1201 Eastlake Avenue East
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98102
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FASTSEQ for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/072,384
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Parker, Gary E
; REGISTRATION NUMBER: 31,648
; REFERENCE/DOCKET NUMBER: 97-16C1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-442-6673
; TELEFAX: 206-442-6678
; TELEX:
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 392 amino acids
; TYPE: amino acid

STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
FRAGMENT TYPE: internal
FEATURE:
NAME/KEY: Signal Sequence
LOCATION: 1...19
OTHER INFORMATION:
US-09-072-384-2

Query Match 97.8%; Score 2085; DB 3; Length 392;
Best Local Similarity 98.2%; Pred. No. 1.8e-220;
Matches 385; Conservative 1; Mismatches 6; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPQCHKGTPPTVEEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRS DGS 120
DB 61 VSSSCGPQCHKGTPPTVEEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRS DGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEKHVLTAACHIDG 180
DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEKHVLTAACHIDG 180

QY 181 KTVVKGTKLRVGLFKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTHTVPKGIKNAND 240
DB 181 KTVVKGTKLRVGLFKPKFKDGGGRANDSTSAMPEQMKFQWIRVKTHTVPKGIKNAND 240

QY 241 IGMDDYDYLLELKKPKFKMKGVSPPAKQLPGGRIHFSGYNDPRGNLVYRFCVDKDE 300
DB 241 IGMDDYDYLLELKKPKFKMKGVSPPAKQLPGGRIHFSGYNDPRGNLVYRFCVDKDE 300

QY 301 TYDLLYQCCDAQPGASGSGVYVVMKRWKQKWKIIGIFSGHQWDMNGSPQDFNVAVR 360
DB 301 TYDLLYQCCDAQPGASGSGVYVVMKRWKQKWKIIGIFSGHQWDMNGSPQDFNVAVR 360

QY 361 ITPLKYAQICYWKGNLYDCREGDTVFPPGNS 392
DB 361 ITPLKYAQICYWKGNLYDCREGDTVFPPGNS 392

RESULT 3

US-09-072-384-18
; Sequence 18, Application US/09072384
; Patent No. 6153420
; GENERAL INFORMATION:
; APPLICANT: Sheppard, Paul O.
; TITLE OF INVENTION: SERINE PROTEASE POLYPEPTIDES
; TITLE OF INVENTION: AND MATERIALS AND METHODS FOR MAKING THEM
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: ZymoGenetics, Inc.
; STREET: 1201 Eastlake Avenue East
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98102
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FASTSEQ for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/072,384
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Parker, Gary E

REGISTRATION NUMBER: 31,648
 REFERENCE/DOCKET NUMBER: 97-16C1
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 206-442-6673
 TELEFAX: 206-442-6678
 TELEX:
 INFORMATION FOR SEQ ID NO: 18:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 383 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 FRAGMENT TYPE: internal
 FEATURE:
 NAME/KEY: Signal Sequence
 LOCATION: 1...19
 OTHER INFORMATION:
 US-09-072-384-18

Query Match 97.6%; Score 2080; DB 3; Length 383;
 Best Local Similarity 100.0%; Pred. No. 6e-220;
 Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MAGIPGLFLFLFLLCAVGVSPYSAPWKPPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
 DB 1 MAGIPGLFLFLFLLCAVGVSPYSAPWKPPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
 QY 61 VSSSCGPOCHKGTPLPTYEAKQYLSYETLYANGSRSTETQVGIYILSSGSGAQHRDGS 120
 DB 61 VSSSCGPOCHKGTPLPTYEAKQYLSYETLYANGSRSTETQVGIYILSSGSGAQHRDGS 120
 QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
 DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
 QY 181 KTVYVGTQKLRVGLFKPKFDGGRGANDSTAMPQMKFQWIRKRVTHVPKGIKGNAND 240
 DB 181 KTVYVGTQKLRVGLFKPKFDGGRGANDSTAMPQMKFQWIRKRVTHVPKGIKGNAND 240
 QY 241 IGMDDYVALLLEKKPKRKFPMKIGVSPPAKQLPGRIHFSGYDNDPRGNLVYRFCDVKDE 300
 DB 241 IGMDDYVALLLEKKPKRKFPMKIGVSPPAKQLPGRIHFSGYDNDPRGNLVYRFCDVKDE 300
 QY 301 TYDLLYQQCDAQPGASGSGVYVWVKRQKQKWKRIIGFSGHWDVMDNGSPQDFNVAVR 360
 DB 301 TYDLLYQQCDAQPGASGSGVYVWVKRQKQKWKRIIGFSGHWDVMDNGSPQDFNVAVR 360
 QY 361 ITPLKVAQICWIKGNVLDREG 383
 DB 361 ITPLKVAQICWIKGNVLDREG 383

RESULT 4
 US-09-907-794A-261
 Sequence 261, Application US/09907794A
 Patent No. 6635468
 GENERAL INFORMATION:
 APPLICANT: Genentech, Inc.
 APPLICANT: Ashkenazi, Avi
 APPLICANT: Botstein, David
 APPLICANT: Desnoyers, Luc
 APPLICANT: Eaton, Dan L.
 APPLICANT: Ferrara, Napoleone
 APPLICANT: Filvaroff, Ellen
 APPLICANT: Fong, Sherman
 APPLICANT: Gao, Wei-Qiang
 APPLICANT: Gerber, Hanspeter
 APPLICANT: Grittisen, Mary E.
 APPLICANT: Goddard, A.
 APPLICANT: Godowski, Paul J.
 APPLICANT: Grimaldi, Christopher J.
 APPLICANT: Gurney, Austin L.

APPLICANT: Hillan, Kenneth, J..
 APPLICANT: Kljavin, Ivar J..
 APPLICANT: Mather, Jemie P..
 APPLICANT: Pan, James
 APPLICANT: Paoni, Nicholas F..
 APPLICANT: Roy, Margaret Ann
 APPLICANT: Stewart, Timothy A..
 APPLICANT: Tumas, Daniel
 APPLICANT: Williams, P. Mickey
 APPLICANT: Wood, William, I..
 TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
 TITLE OF INVENTION: Acids Encoding the Same
 FILE REFERENCE: 10466-14
 CURRENT APPLICATION NUMBER: US/09/907,794A
 CURRENT FILING DATE: 2001-07-17
 PRIOR APPLICATION NUMBER: PCT/US00/04414
 PRIOR FILING DATE: 2000-02-22
 PRIOR APPLICATION NUMBER: US 60/143,048
 PRIOR FILING DATE: 1999-07-07
 PRIOR APPLICATION NUMBER: US 60/145,698
 PRIOR FILING DATE: 1999-07-26
 PRIOR APPLICATION NUMBER: US 60/146,222
 PRIOR FILING DATE: 1999-07-28
 PRIOR APPLICATION NUMBER: PCT/US99/20594
 PRIOR FILING DATE: 1999-09-08
 PRIOR APPLICATION NUMBER: PCT/US99/20944
 PRIOR FILING DATE: 1999-09-13
 PRIOR APPLICATION NUMBER: PCT/US99/21090
 PRIOR FILING DATE: 1999-09-15
 PRIOR APPLICATION NUMBER: PCT/US99/21547
 PRIOR FILING DATE: 1999-09-15
 PRIOR APPLICATION NUMBER: PCT/US99/23089
 PRIOR FILING DATE: 1999-10-05
 PRIOR APPLICATION NUMBER: PCT/US99/28214
 PRIOR FILING DATE: 1999-11-29
 PRIOR APPLICATION NUMBER: PCT/US99/28313
 PRIOR FILING DATE: 1999-11-30
 PRIOR APPLICATION NUMBER: PCT/US99/28564
 PRIOR FILING DATE: 1999-12-02
 PRIOR APPLICATION NUMBER: PCT/US99/28565
 PRIOR FILING DATE: 1999-12-02
 PRIOR APPLICATION NUMBER: PCT/US99/30095
 PRIOR FILING DATE: 1999-12-16
 PRIOR APPLICATION NUMBER: PCT/US99/30911
 PRIOR FILING DATE: 1999-12-20
 PRIOR APPLICATION NUMBER: PCT/US99/30999
 PRIOR FILING DATE: 1999-12-20
 PRIOR APPLICATION NUMBER: PCT/US00/00219
 PRIOR FILING DATE: 2000-01-05
 NUMBER OF SEQ ID NOS: 423
 SEQ ID NO 261
 LENGTH: 383
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-09-907-794A-261
 Query Match 97.6%; Score 2080; DB 4; Length 383;
 Best Local Similarity 100.0%; Pred. No. 6e-220;
 Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MAGIPGLFLFLFLLCAVGVSPYSAPWKPPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
 DB 1 MAGIPGLFLFLFLLCAVGVSPYSAPWKPPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
 QY 61 VSSSCGPOCHKGTPLPTYEAKQYLSYETLYANGSRSTETQVGIYILSSGSGAQHRDGS 120
 DB 61 VSSSCGPOCHKGTPLPTYEAKQYLSYETLYANGSRSTETQVGIYILSSGSGAQHRDGS 120
 QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
 DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
 QY 181 KTVYVGTQKLRVGLFKPKFDGGRGANDSTAMPQMKFQWIRKRVTHVPKGIKGNAND 240

Db 181 KTYVKGQKLRVGLPKPKFKDGGGRANDSTSAPEQMKFQWIRVKTHTVPGKWIKNAND 240
Qy 241 IGMDDYALLELKKPKRKFPMKIGVSPPAKQPGRIHFSGYDNDPRPGLVYRFCVDVKDE 300
Db 241 IGMDDYALLELKKPKRKFPMKIGVSPPAKQPGRIHFSGYDNDPRPGLVYRFCVDVKDE 300
Qy 301 TYDLLYQQCDAQPGASGGVYVRMWKQQQKWERKIIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAQPGASGGVYVRMWKQQQKWERKIIIGIFSGHQWDMNGSPQDFNVAVR 360
Qy 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 5
US-09-905-125A-261
; Sequence 261, Application US/09905125A
; Patent No. 6664376
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,125A
; PRIOR FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30

; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-905-125A-261
Query Match 97.6%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 6e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 MAGIFGLLELFLFLICAVGQVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKPDPGAEAKLE 60
Db 1 MAGIFGLLELFLFLICAVGQVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKPDPGAEAKLE 60
Qy 61 VSSSCGPOCHKGTPLPTTVEEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDSSG 120
Db 61 VSSSCGPOCHKGTPLPTTVEEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDSSG 120
Qy 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTCTGLVAEKHVLTAACHIDG 180
Qy 181 KTYVKGQKLRVGLPKPKFKDGGGRANDSTSAPEQMKFQWIRVKTHTVPGKWIKNAND 240
Db 181 KTYVKGQKLRVGLPKPKFKDGGGRANDSTSAPEQMKFQWIRVKTHTVPGKWIKNAND 240
Qy 241 IGMDDYALLELKKPKRKFPMKIGVSPPAKQPGRIHFSGYDNDPRPGLVYRFCVDVKDE 300
Db 241 IGMDDYALLELKKPKRKFPMKIGVSPPAKQPGRIHFSGYDNDPRPGLVYRFCVDVKDE 300
Qy 301 TYDLLYQQCDAQPGASGGVYVRMWKQQQKWERKIIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAQPGASGGVYVRMWKQQQKWERKIIIGIFSGHQWDMNGSPQDFNVAVR 360
Qy 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 6
US-09-902-775A-261
; Sequence 261, Application US/09902775A
; Patent No. 6686451
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.

APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/902,775A
PRIOR FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/30095
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-902-775A-261

Query Match 97.6%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 68-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAGGVSPYAPWKPWPAYRLPVVLPSTLNLAQPDFAEAKLE 60
DB 1 MAGIPGLLFLFLLCAGGVSPYAPWKPWPAYRLPVVLPSTLNLAQPDFAEAKLE 60
QY 61 VSSSCGPGQCHKGTPLTYEAKQVLSVETLYANGSRRTQVGIYIILSSSGDGAQHRDGS 120
DB 61 VSSSCGPGQCHKGTPLTYEAKQVLSVETLYANGSRRTQVGIYIILSSSGDGAQHRDGS 120
QY 121 SGKSRKRQIYGVDSRFSIFKDFLLNYPSTSVKLSGTCTGLTVAEKHVLTAACHIDG 180
DB 121 SGKSRKRQIYGVDSRFSIFKDFLLNYPSTSVKLSGTCTGLTVAEKHVLTAACHIDG 180
QY 181 KTVVKGTKLVRGFLKPKFGDGRANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240

DB 181 KTVVKGTKLVRGFLKPKFGDGRANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
QY 241 IGMDYDYALLELKPKHKRKFMIKIGVSPPAKQIPGGRIHESGYDNDPBGNLVYRFFCDVKDE 300
DB 241 IGMDYDYALLELKPKHKRKFMIKIGVSPPAKQIPGGRIHESGYDNDPBGNLVYRFFCDVKDE 300
QY 301 TYDLLYQCDQAQGASGSGVYVMMKRCQQKWKRIIGIFSGHQWYDMNGSPQDFNVAVR 360
DB 301 TYDLLYQCDQAQGASGSGVYVMMKRCQQKWKRIIGIFSGHQWYDMNGSPQDFNVAVR 360
QY 361 ITPLKYAQICYWIKGNLYDCREG 383
DB 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 7
US-09-906-700-261
Sequence 261, Application US/09906700
Patent No. 6723535
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/906,700
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564

; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-906-700-261

Query Match 97.6%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 6e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPPTWPAYRLPVVLPQSTLNLAKPDPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPPTWPAYRLPVVLPQSTLNLAKPDPFGAEAKLE 60

Qy 61 VSSSCGPOCHKGTPLPTYEAAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDSDS 120
Db 61 VSSSCGPOCHKGTPLPTYEAAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDSDS 120

Qy 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180

Qy 181 KTYVKGTKLRVGFLLKPKFKDGGGRANDSTSAMPQMKFQWIRVKRTHVTPKGIKGNAND 240
Db 181 KTYVKGTKLRVGFLLKPKFKDGGGRANDSTSAMPQMKFQWIRVKRTHVTPKGIKGNAND 240

Qy 241 IGMDDYALLELKKPHKRFMKIGVSPPAKQLPGGRIFHSFGYDNDPRGNLVYRFCDDVDE 300
Db 241 IGMDDYALLELKKPHKRFMKIGVSPPAKQLPGGRIFHSFGYDNDPRGNLVYRFCDDVDE 300

Qy 301 TYDLLVQCCDAQPGASGSGVYVVMWKRQQQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLVQCCDAQPGASGSGVYVVMWKRQQQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360

Qy 361 ITPKYAQICYWKIGNYLDREG 383
Db 361 ITPKYAQICYWKIGNYLDREG 383

RESULT 8
US-09-903-603A-261
; Sequence 261, Application US/09903603A
; Patent No. 6767995
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Aabkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.

; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: GNE.1618P2C12
; CURRENT APPLICATION NUMBER: US/09/903,603A
; CURRENT FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-903-603A-261

Query Match 97.6%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 6e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPPTWPAYRLPVVLPQSTLNLAKPDPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPPTWPAYRLPVVLPQSTLNLAKPDPFGAEAKLE 60

Qy 61 VSSSCGPOCHKGTPLPTYEAAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDSDS 120
Db 61 VSSSCGPOCHKGTPLPTYEAAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDSDS 120

Qy 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180

Qy 181 KTYVKGTKLRVGFLLKPKFKDGGGRANDSTSAMPQMKFQWIRVKRTHVTPKGIKGNAND 240
Db 181 KTYVKGTKLRVGFLLKPKFKDGGGRANDSTSAMPQMKFQWIRVKRTHVTPKGIKGNAND 240

Qy	241	IGWDYDYALLELKKPHKKFKMKI	GVSPPAKQL	PGGRTH	FSGYDND	RPGNLVY	RFCDVKDE	300
Db	241	IGWDYDYALLELKKPHKKFKMKI	GVSPPAKQL	PGGRTH	FSGYDND	RPGNLVY	RFCDVKDE	300
Qy	301	TYDLLYQQCDAQ	PGAGSGSVV	VYRWKROQ	QKWERKII	GIFSGHQW	DMNGSPQDFNVAVR	360
Db	301	TYDLLYQQCDAQ	PGAGSGSVV	VYRWKROQ	QKWERKII	GIFSGHQW	DMNGSPQDFNVAVR	360
Qy	361	ITPLKYAQIC	YWKIGNYLD	CREG	383			
Db	361	ITPLKYAQIC	YWKIGNYLD	CREG	383			

RESULT 9

US-09-904-920A-261
 ; Sequence 261, Application US/09904920A
 ; Patent No. 6806352
 ; GENERAL INFORMATION:
 ; APPLICANT: Genentech, Inc.
 ; APPLICANT: Ashkenazi, Avi
 ; APPLICANT: Botstein, David
 ; APPLICANT: Desnoyers, Luc
 ; APPLICANT: Eaton, Dan L.
 ; APPLICANT: Ferrara, Napoleone
 ; APPLICANT: Filvaroff, Ellen
 ; APPLICANT: Fong, Sherman
 ; APPLICANT: Gao, Wei-Qiang
 ; APPLICANT: Gerber, Hanspeter
 ; APPLICANT: Gerritsen, Mary E.
 ; APPLICANT: Goddard, A.
 ; APPLICANT: Godowski, Paul J.
 ; APPLICANT: Grimaldi, Christopher J.
 ; APPLICANT: Gurney, Austin L.
 ; APPLICANT: Hillan, Kenneth, J.
 ; APPLICANT: Kijavlin, Ivar J.
 ; APPLICANT: Mather, Jennie P.
 ; APPLICANT: Pan, James
 ; APPLICANT: Paoni, Nicholas F.
 ; APPLICANT: Roy, Margaret Ann
 ; APPLICANT: Stewart, Timothy A.
 ; APPLICANT: Tumas, Daniel
 ; APPLICANT: Williams, P. Mickey
 ; APPLICANT: Wood, William, I.
 ; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
 ; TITLE OF INVENTION: Acids Encoding the Same
 ; FILE REFERENCE: 10466-14
 ; CURRENT APPLICATION NUMBER: US/09/904,920A
 ; CURRENT FILING DATE: 2001-07-13
 ; PRIOR APPLICATION NUMBER: PCT/US00/04414
 ; PRIOR FILING DATE: 2000-02-22
 ; PRIOR APPLICATION NUMBER: US 60/143,048
 ; PRIOR FILING DATE: 1999-07-07
 ; PRIOR APPLICATION NUMBER: US 60/145,698
 ; PRIOR FILING DATE: 1999-07-26
 ; PRIOR APPLICATION NUMBER: US 60/146,222
 ; PRIOR FILING DATE: 1999-07-28
 ; PRIOR APPLICATION NUMBER: PCT/US99/20594
 ; PRIOR FILING DATE: 1999-09-08
 ; PRIOR APPLICATION NUMBER: PCT/US99/20944
 ; PRIOR FILING DATE: 1999-09-13
 ; PRIOR APPLICATION NUMBER: PCT/US99/21090
 ; PRIOR FILING DATE: 1999-09-15
 ; PRIOR APPLICATION NUMBER: PCT/US99/21547
 ; PRIOR FILING DATE: 1999-09-15
 ; PRIOR APPLICATION NUMBER: PCT/US99/23089
 ; PRIOR FILING DATE: 1999-10-05
 ; PRIOR APPLICATION NUMBER: PCT/US99/28214
 ; PRIOR FILING DATE: 1999-11-29
 ; PRIOR APPLICATION NUMBER: PCT/US99/28313
 ; PRIOR FILING DATE: 1999-11-30
 ; PRIOR APPLICATION NUMBER: PCT/US99/28564
 ; PRIOR FILING DATE: 1999-12-02

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; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-304-920A-261

Query Match          97.6%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 6e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFLLLCAVGQSPYSPAPWKPTWPAYRLPVVLPQSTLNLA KP DFGAEAKLE 60
Db 1 MAGIPGLLFLFLLLCAVGQSPYSPAPWKPTWPAYRLPVVLPQSTLNLA KP DFGAEAKLE 60
Qy 61 VSSSCGPGQCHKGTPLPTYBEAKQYLSYETIYANGSRRTETQVGIYILSSSGDGAQHRDSGS 120
Db 61 VSSSCGPGQCHKGTPLPTYBEAKQYLSYETIYANGSRRTETQVGIYILSSSGDGAQHRDSGS 120
Qy 121 SGKRRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTGCTLVAEKHVLTAACHIDG 180
Db 121 SGKRRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTGCTLVAEKHVLTAACHIDG 180
Qy 181 KTYVGTQKLRVGLFKPKFQDGGRGANDSTSAMPEQMKFQWIRKRTHPKGIWIKGNAND 240
Db 181 KTYVGTQKLRVGLFKPKFQDGGRGANDSTSAMPEQMKFQWIRKRTHPKGIWIKGNAND 240
Qy 241 IGMDDYALLELKKPHKRKPMKI GVSPPAKQLPGGRIFHFGSYNDRPGLNLYRFCDVKDE 300
Db 241 IGMDDYALLELKKPHKRKPMKI GVSPPAKQLPGGRIFHFGSYNDRPGLNLYRFCDVKDE 300
Qy 301 TYDLLVQQCDAQPGASGSGYVVRWKRQOQKWERKIIGFSGHOWDMNGSPQDFNVAVR 360
Db 301 TYDLLVQQCDAQPGASGSGYVVRWKRQOQKWERKIIGFSGHOWDMNGSPQDFNVAVR 360
Qy 361 ITELKVAQICYIKGNLYDCREG 383
Db 361 ITELKVAQICYIKGNLYDCREG 383

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RESULT 10

US-09-309-064-261
; Patent 261, Application US/09309064
; Patent No. 6819449
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Bocstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavlin, Ivar J.
; APPLICANT: Mather, Jennie P.

```

; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/909,064
; PRIOR FILING DATE: 2001-07-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-909-064-261

Query Match          97.6%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 6e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1  MAGIPGLLFLFLLCVAGQVSPYAPWKPWPAYRLPVVLPSTLNLAQPDFAEAKLE 60
DB      1  MAGIPGLLFLFLLCVAGQVSPYAPWKPWPAYRLPVVLPSTLNLAQPDFAEAKLE 60

QY      61  VSSSCGPQCHKGTPLPYEEAKQYLSYETLYANGSRRTQVGYIILSSSGDGAQHRDGS 120
DB      61  VSSSCGPQCHKGTPLPYEEAKQYLSYETLYANGSRRTQVGYIILSSSGDGAQHRDGS 120

QY      121  SGKSRKRQIYGVDSRFSFGKDFLNNPSTSVKLSCTGCTGLVAEKHVLTAACHIDG 180
DB      121  SGKSRKRQIYGVDSRFSFGKDFLNNPSTSVKLSCTGCTGLVAEKHVLTAACHIDG 180

QY      181  KTVVKGQTQLRVGFLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240
DB      181  KTVVKGQTQLRVGFLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240
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QY      241  IGMVDYDVALLELKPKHKKFMKIGVSPRAKQLPGGRIHESGYDNDPRPGNLVYRFCVDKDE 300
DB      241  IGMVDYDVALLELKPKHKKFMKIGVSPRAKQLPGGRIHESGYDNDPRPGNLVYRFCVDKDE 300

QY      301  TYDLLYQQCDAQFAGSGSGVYVVMWKRQOQKWERKIIGIFSGHQMVMNGSPQDFNVAVR 360
DB      301  TYDLLYQQCDAQFAGSGSGVYVVMWKRQOQKWERKIIGIFSGHQMVMNGSPQDFNVAVR 360

QY      361  ITPLKYAQICYWIKGNLYLDCREG 383
DB      361  ITPLKYAQICYWIKGNLYLDCREG 383

RESULT 11
US-09-905-381A-261
; Sequence 261, Application US/09905381A
; Patent No. 6818746
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,381A
; PRIOR FILING DATE: 2001-07-13
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
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; PRIOR APPLICATION NUMBER: PCT/US99/28214
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; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
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; PRIOR FILING DATE: 1999-12-02
; -PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-905-381A-261

Query March 97.6%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 6e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0

Qy 1 MAGIPGGLLFLFFLLCAVGQSPSPAPKPTWPAVRLPVLPQSTLNIAKDFGAEAKLE 60
Db 1 MAGIPGGLLFLFFLLCAVGQSPSPAPKPTWPAVRLPVLPQSTLNIAKDFGAEAKLE 60

Qy 61 VSSSCGPOCHKGTPLPTVEEAKQVLSYETLVANGSRSTQVGIYILSSSGDGAQRDSSG 120
Db 61 VSSSCGPOCHKGTPLPTVEEAKQVLSYETLVANGSRSTQVGIYILSSSGDGAQRDSSG 120

Qy 121 SGKRRKRQIYGYDSRSPISFKOKFLNYPFSTSVKLSCTGCTGLVAEKHLVTAACHIDG 180
Db 121 SGKRRKRQIYGYDSRSPISFKOKFLNYPFSTSVKLSCTGCTGLVAEKHLVTAACHIDG 180

Qy 181 KTYVYGTQKLRVGLPKPKFDGGRGANDSTSAMPEOMKFQWIRVKRTHVPKGIKGNAND 240
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Qy 241 IGMDDYDALLLEKKPHKRKFMKIIGVSPPAKOLPGRHFSGVDNDRPGNLVYRFDVKDE 300
Db 241 IGMDDYDALLLEKKPHKRKFMKIIGVSPPAKOLPGRHFSGVDNDRPGNLVYRFDVKDE 300

Qy 301 TYDLLYQQCDAQPCGASGSGVYVRMWKRRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAQPCGASGSGVYVRMWKRRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360

Qy 361 ITPLKYAQICWIKGNLYDCREG 383
Db 361 ITPLKYAQICWIKGNLYDCREG 383

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RESULT 12

US-09-906-618-261

Sequence 261, Application US/09906618

Patent No. 6828146

GENERAL INFORMATION:

APPLICANT: Genentech, Inc.

APPLICANT: Ashkenazi, Avi

APPLICANT: Botstein, David

APPLICANT: Deonoyers, Luc

APPLICANT: Eaton, Dan L.

APPLICANT: Ferrara, Napoleone

APPLICANT: Filvaroff, Ellen

APPLICANT: Fong, Sherman

APPLICANT: Gao, Wei-Qiang

APPLICANT: Gerber, Hanspeter

APPLICANT: Gerritsen, Mary E.

APPLICANT: Goddard, A.

APPLICANT: Godowski, Paul J.

APPLICANT: Grimaldi, Christopher J.

APPLICANT: Gurney, Austin L.

APPLICANT: Hillan, Kenneth, J.

APPLICANT: Kijavlin, Ivar J.

APPLICANT: Mather, Jennie P.

APPLICANT: Pan, James

```

1  APPLICANT: Paoni, Nicholas F.
2  APPLICANT: Roy, Margaret Ann
3  APPLICANT: Stewart, Timothy A.
4  APPLICANT: Tumas, Daniel
5  APPLICANT: Williams, P. Mickey
6  APPLICANT: Wood, William, I.
7  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
8  TITLE OF INVENTION: Acids Encoding the Same
9  FILE REFERENCE: 10466-14
10 CURRENT APPLICATION NUMBER: US/09/906.618
11 CURRENT FILING DATE: 2001-07-16
12 PRIOR APPLICATION NUMBER: PCT/US00/04414
13 PRIOR FILING DATE: 2000-02-22
14 PRIOR APPLICATION NUMBER: US 60/143,048
15 PRIOR FILING DATE: 1999-07-07
16 PRIOR APPLICATION NUMBER: US 60/145,698
17 PRIOR FILING DATE: 1999-07-26
18 PRIOR APPLICATION NUMBER: US 60/146,222
19 PRIOR FILING DATE: 1999-07-28
20 PRIOR APPLICATION NUMBER: PCT/US99/20594
21 PRIOR FILING DATE: 1999-09-08
22 PRIOR APPLICATION NUMBER: PCT/US99/20944
23 PRIOR FILING DATE: 1999-09-13
24 PRIOR APPLICATION NUMBER: PCT/US99/21090
25 PRIOR FILING DATE: 1999-09-15
26 PRIOR APPLICATION NUMBER: PCT/US99/21547
27 PRIOR FILING DATE: 1999-09-15
28 PRIOR APPLICATION NUMBER: PCT/US99/23089
29 PRIOR FILING DATE: 1999-10-05
30 PRIOR APPLICATION NUMBER: PCT/US99/28214
31 PRIOR FILING DATE: 1999-11-29
32 PRIOR APPLICATION NUMBER: PCT/US99/28313
33 PRIOR FILING DATE: 1999-11-30
34 PRIOR APPLICATION NUMBER: PCT/US99/28564
35 PRIOR FILING DATE: 1999-12-02
36 PRIOR APPLICATION NUMBER: PCT/US99/28565
37 PRIOR FILING DATE: 1999-12-02
38 PRIOR APPLICATION NUMBER: PCT/US99/30095
39 PRIOR FILING DATE: 1999-12-16
40 PRIOR APPLICATION NUMBER: PCT/US99/30911
41 PRIOR FILING DATE: 1999-12-20
42 PRIOR APPLICATION NUMBER: PCT/US99/30999
43 PRIOR FILING DATE: 1999-12-20
44 PRIOR APPLICATION NUMBER: PCT/US00/00219
45 PRIOR FILING DATE: 2000-01-05
46 NUMBER OF SEQ ID NOS: 423
47 SEQ ID NO 261
48 LENGTH: 383
49 TYPE: PRT
50 ORGANISM: Homo Sapien
51 US-09-906-618-261

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Db	121	SGKSRKRQIYGYDYSRFSIFGKDF	FLNYPSTSVKLS	TGCTGTLVA	SKHVLTA	AHC1HDG 180	
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Db 361 ITPKVAQICYWIKGNLYDCREG 383

RESULT 13
US-09-551-826D-6
; Sequence 6, Application US/09551826D.
; Patent No. 6558939
; GENERAL INFORMATION:
; APPLICANT: No. 6558939regaard-Madsen, Mads
; APPLICANT: Ostergaard, Peter Rahbek
; APPLICANT: Christensen, Claus Bo Voge
; APPLICANT: Lassen, Soren Plensted
; TITLE OF INVENTION: No. 6558939el Poteases And Variants Thereof
; FILE REFERENCE: 5665.200-US
; CURRENT APPLICATION NUMBER: US/09/551,826D
; CURRENT FILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 6
; LENGTH: 314
; TYPE: PRT
; ORGANISM: Bacillus licheniformis AC116
US-09-551-826D-6

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Matches 77; Conservative 37; Mismatches 137; Indels 59; Gaps 13;

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Qy 133 YDSRFSIFGKDFLLNPFSTSVKLTST---GCTGTVAEKHVLTAACIHD-KTYTVKGTQ 188
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Qy 249 LLELKKPKHKKFPMKIGVSPPAKQLPGGRHFGSYDNDPRGNLVRFCDD---VKDETLYDLY 306
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Qy 307 QCCDAQPGASGSGVYVMWKRQOQKWERKII-----GIFSGHQWDMNGSPQDFNVAVRIT 362
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RESULT 14
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; Sequence 2, Application US/09551826D
; Patent No. 6558939
; GENERAL INFORMATION:
; APPLICANT: No. 6558939regaard-Madsen, Mads
; APPLICANT: Ostergaard, Peter Rahbek
; APPLICANT: Christensen, Claus Bo Voge
; APPLICANT: Lassen, Soren Plensted
; TITLE OF INVENTION: No. 6558939el Poteases And Variants Thereof
; FILE REFERENCE: 5665.200-US
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; CURRENT APPLICATION NUMBER: US/09/551,826D
; CURRENT FILING DATE: 2000-04-17
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 316
; TYPE: PRT
; ORGANISM: Bacillus licheniformis
US-09-551-826D-2

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Matches 79; Conservative 43; Mismatches 143; Indels 64; Gaps 15;

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RESULT 15
US-08-090-048-1
; Sequence 1, Application US/080900048
; Patent No. 5523237
; GENERAL INFORMATION:
; APPLICANT: Budtz, Peter
; APPLICANT: Nielsen, Per M.
; TITLE OF INVENTION: PROTEIN PREPARATIONS
; NUMBER OF SEQUENCES: 2
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: No. 5523237o No. 5523237disk of No. 5523237th America, Inc.
; STREET: 405 Lexington Avenue, 62nd Floor
; CITY: New York
; STATE: New York
; COUNTRY: United States of America
; ZIP: 10174-6201
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/090,048
; FILING DATE: 16-JUL-1993
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: DK 199/91
; FILING DATE: 06-FEB-1991
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: PCT/DK92/00036
; FILING DATE: 06-FEB-1992
; ATTORNEY/AGENT INFORMATION:
; NAME: Agriis, Cheryl H.
; REGISTRATION NUMBER: 34,086
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; REFERENCE/DOCKET NUMBER: 3396.214-US
 ; TELECOMMUNICATION INFORMATION:
 ; TELEPHONE: 212-867-0123
 ; TELEFAX: 212-867-0298
 ; INFORMATION FOR SEQ ID NO: 1:
 ; SEQUENCE CHARACTERISTICS:
 ; LENGTH: 222 amino acids
 ; TYPE: amino acid
 ; STRANDEDNESS: single
 ; TOPOLOGY: linear
 ; MOLECULE TYPE: protein
 ; US-08-090-048-1

Query Match 8.3%; Score 177; DB 1; Length 222;
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DB	67	---	RNGTS	----	YPYGSVKSTR	YFIPSGWRSGNTN	-----	113	
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

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(without alignments)
1673.692 Million cell updates/sec

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Minimum DB seq length: 0
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Maximum Match 100%
Listing first 45 summaries

Database : Published Applications AA:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	2080	97.6	383	9	US-09-765-205-12
2	2080	97.6	383	9	US-09-909-320-261
3	2080	97.6	383	9	US-09-909-088B-261
4	2080	97.6	383	9	US-09-905-291A-261
5	2080	97.6	383	9	US-09-902-853-261
6	2080	97.6	383	9	US-09-907-824-261
7	2080	97.6	383	9	US-09-907-841-261
8	2080	97.6	383	10	US-09-904-011-261
9	2080	97.6	383	10	US-09-903-640-261
10	2080	97.6	383	10	US-09-908-093-261
11	2080	97.6	383	10	US-09-906-742-261

12	2080	97.6	383	10	US-09-906-838-261	Sequence 261, App
13	2080	97.6	383	10	US-09-907-613-261	Sequence 261, App
14	2080	97.6	383	10	US-09-907-942-261	Sequence 261, App
15	2080	97.6	383	10	US-09-904-859-261	Sequence 261, App
16	2080	97.6	383	10	US-09-903-204-261	Sequence 261, App
17	2080	97.6	383	10	US-09-904-820-261	Sequence 261, App
18	2080	97.6	383	10	US-09-904-786-261	Sequence 261, App
19	2080	97.6	383	10	US-09-906-646-261	Sequence 261, App
20	2080	97.6	383	10	US-09-906-700-261	Sequence 261, App
21	2080	97.6	383	10	US-09-903-786-261	Sequence 261, App
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24	2080	97.6	383	10	US-09-904-119-261	Sequence 261, App
25	2080	97.6	383	10	US-09-904-956-261	Sequence 261, App
26	2080	97.6	383	10	US-09-902-736-261	Sequence 261, App
27	2080	97.6	383	10	US-09-907-794-261	Sequence 261, App
28	2080	97.6	383	10	US-09-903-943-261	Sequence 261, App
29	2080	97.6	383	10	US-09-904-462-261	Sequence 261, App
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31	2080	97.6	383	10	US-09-902-692-261	Sequence 261, App
32	2080	97.6	383	10	US-09-903-520-261	Sequence 261, App
33	2080	97.6	383	10	US-09-905-056-261	Sequence 261, App
34	2080	97.6	383	10	US-09-984-130-45	Sequence 45, Appl
35	2080	97.6	383	10	US-09-984-130-126	Sequence 126, App
36	2080	97.6	383	10	US-09-909-064-261	Sequence 261, App
37	2080	97.6	383	10	US-09-904-553-261	Sequence 261, App
38	2080	97.6	383	10	US-09-905-381-261	Sequence 261, App
39	2080	97.6	383	10	US-09-904-485-261	Sequence 261, App
40	2080	97.6	383	10	US-09-905-348-261	Sequence 261, App
41	2080	97.6	383	10	US-09-905-088-261	Sequence 261, App
42	2080	97.6	383	10	US-09-907-575-261	Sequence 261, App
43	2080	97.6	383	10	US-09-905-075-261	Sequence 261, App
44	2080	97.6	383	10	US-09-902-759-261	Sequence 261, App
45	2080	97.6	383	10	US-09-902-634-261	Sequence 261, App

ALIGNMENTS

RESULT 1
US-09-765-205-12
; Sequence 12, Application US/09765205
; Patent No. US20020034800A1
; GENERAL INFORMATION:
; APPLICANT: Cao, Li
; TITLE OF INVENTION: BONE MARROW SECRETED PROTEINS AND POLYNUCLEOTIDES
; FILE REFERENCE: 1458.004/200130.449
; CURRENT APPLICATION NUMBER: US/09/765,205
; CURRENT FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: US/09/212,440
; PRIOR FILING DATE: 1998-12-16
; NUMBER OF SEQ ID NOS: 46
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 12
; LENGTH: 383
; TYPE: PRT
; ORGANISM: human
; US-09-765-205-12

Query Match		97.6%	Score 2080;	DB 9;	Length 383;
Best Local Similarity		100.0%;	Pred. No. 2,2e-193;		
Matches 383;		Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
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DB	61	VSSSCGPGQCHKGTPTPTTBEAKQYLSYETLYANGRTETQVGIYI	LSSSGDGA	QHRD	SGS 120
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Db 301 TYDLLYQCCDAQPGASGSGVYVVMWKROQKWKRIIGIFSGHGWDMNGSPQDFNVAVR 360
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RESULT 2
US-09-909-320-261
; Sequence 261, Application US/09909320
; Patent No. US20020132240A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/909,320
; PRIOR FILING DATE: 2002-01-04
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
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; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-909-320-261

Query Match 97.6%; Score 2080; DB 9; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.2e-193; Mismatches 0; Indels 0; Gaps 0;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFFLLCAVGVQSPYSAPWKPWPAYRLPVLPQSTINLAKPDEGAEAKLE 60
Db 1 MAGIPGLLFLFFLLCAVGVQSPYSAPWKPWPAYRLPVLPQSTINLAKPDEGAEAKLE 60
QY 61 VSSSCGPGCHKGTPLPYEEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDGS 120
Db 61 VSSSCGPGCHKGTPLPYEEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDGS 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
QY 181 KTYVKGTKLRVGLFKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
Db 181 KTYVKGTKLRVGLFKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
QY 241 IGMDDYDYLLELKKPHKPKFKMGKIGVSPPAKQLPGGRIFHSGYDNDPRGNLVRFCVDKDE 300
Db 241 IGMDDYDYLLELKKPHKPKFKMGKIGVSPPAKQLPGGRIFHSGYDNDPRGNLVRFCVDKDE 300
QY 301 TYDLLYQCCDAQPGASGSGVYVVMWKROQKWKRIIGIFSGHGWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAQPGASGSGVYVVMWKROQKWKRIIGIFSGHGWDMNGSPQDFNVAVR 360
QY 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 3
US-09-909-088B-261
; Sequence 261, Application US/09909088B
; Patent No. US20020146709A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.

```

; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/909,088B
; PRIOR FILING DATE: 2001-07-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-909-088B-261

Query Match          97.6%; Score 2080; DB 9; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.2e-193;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLPFLCAVGQVSPYAPWKTWPAYRLPVVLPSTLNLAQPDGAEAKLE 60
Db 1 MAGIPGLLFLPFLCAVGQVSPYAPWKTWPAYRLPVVLPSTLNLAQPDGAEAKLE 60
Qy 61 VSSSCGQCHKGTPLTYEAKQYLSYETLYANGSRTQVGIYILSSGSDGAQHRDSCS 120
Db 61 VSSSCGQCHKGTPLTYEAKQYLSYETLYANGSRTQVGIYILSSGSDGAQHRDSCS 120
Qy 121 SGKSRKRQIYGYDSRFSFGKDFLNLNYPSTSVKLSGTCTGLVAEKHVLTAACHIHG 180
Db 121 SGKSRKRQIYGYDSRFSFGKDFLNLNYPSTSVKLSGTCTGLVAEKHVLTAACHIHG 180
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Qy 181 KTYVKGTKLRVGFGLKPKFKGCGRGANDSTSAMPEOMKQWIRVKTHTVPGKWIKNAND 240
Db 181 KTYVKGTKLRVGFGLKPKFKGCGRGANDSTSAMPEOMKQWIRVKTHTVPGKWIKNAND 240
Qy 241 IGMVDYDYLLELLEKPKHKKFKMKGIVSPPAKQLPGGRIHFSGYDNDPRGNLVYRFCDVKDE 300
Db 241 IGMVDYDYLLELLEKPKHKKFKMKGIVSPPAKQLPGGRIHFSGYDNDPRGNLVYRFCDVKDE 300
Qy 301 TYDLLYQCCDAOPGASGSGVYVMKQRQOKKVERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAOPGASGSGVYVMKQRQOKKVERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Qy 361 ITPLKYAQICYWIKGNLYLDCREG 383
Db 361 ITPLKYAQICYWIKGNLYLDCREG 383

RESULT 4
US-09-905-291A-261
; Sequence 261, Application US/09905291A
; Patent No. US20020160374A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,291A
; PRIOR FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
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QY	181	KTYYVGTQKLRVGFLLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND	240
Db	181	KTYYVGTQKLRVGFLLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND	240
QY	241	IGMDYDVALLELKKPKKFKMKIGVSPPAKQLPGRTHFGSYDNDRPGNLVYRFFCDVKDE	300
Db	241	IGMDYDVALLELKKPKKFKMKIGVSPPAKQLPGRTHFGSYDNDRPGNLVYRFFCDVKDE	300
QY	301	TYDLYVQCDAQPGASGSGVYVVMKRWKQQKWKERKIIIGIFSGHQWVDMNGSPQDFNVAVR	360
Db	301	TYDLYVQCDAQPGASGSGVYVVMKRWKQQKWKERKIIIGIFSGHQWVDMNGSPQDFNVAVR	360
QY	361	ITPLKYAQICYWIKGNLYDCREG	383
Db	361	ITPLKYAQICYWIKGNLYDCREG	383

RESULT 6

US-09-907-824-261

Sequence 261, Application US/09907824

Publication No. US20020197671A1

GENERAL INFORMATION:

APPLICANT: Genentech, Inc.

APPLICANT: Ashkenazi, Avi

APPLICANT: Botstein, David

APPLICANT: Desnoyers, Luc

APPLICANT: Eaton, Dan L.

APPLICANT: Ferrara, Napoleone

APPLICANT: Filvaroff, Ellen

APPLICANT: Fong, Sherman

APPLICANT: Gao, Wei-Qiang

APPLICANT: Gerber, Hanspeter

APPLICANT: Gerritsen, Mary E.

APPLICANT: Goddard, A.

APPLICANT: Godowski, Paul J.

APPLICANT: Grimaldi, Christopher J.

APPLICANT: Gurney, Austin L.

APPLICANT: Hillan, Kenneth, J.

APPLICANT: Kljavin, Ivar J.

APPLICANT: Mather, Jennie P.

APPLICANT: Pan, James

APPLICANT: Paoni, Nicholas F.

APPLICANT: Roy, Margaret Ann

APPLICANT: Stewart, Timothy A.

APPLICANT: Tumas, Daniel

APPLICANT: Williams, P. Mickey

APPLICANT: Wood, William, I.

TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic

TITLE OF INVENTION: Acids Encoding the Same

FILE REFERENCE: 10466-14

CURRENT APPLICATION NUMBER: US/09/907,824

PRIOR FILING DATE: 2001-07-17

PRIOR APPLICATION NUMBER: 09/665,350

PRIOR FILING DATE: 2000-09-18

PRIOR APPLICATION NUMBER: PCT/US00/04414

PRIOR FILING DATE: 2000-02-22

PRIOR APPLICATION NUMBER: US 60/143,048

PRIOR FILING DATE: 1999-07-07

PRIOR APPLICATION NUMBER: US 60/145,698

PRIOR FILING DATE: 1999-07-26

PRIOR APPLICATION NUMBER: US 60/146,222

PRIOR FILING DATE: 1999-07-28

PRIOR APPLICATION NUMBER: PCT/US99/20594

PRIOR FILING DATE: 1999-09-08

PRIOR APPLICATION NUMBER: PCT/US99/20944

PRIOR FILING DATE: 1999-09-13

PRIOR APPLICATION NUMBER: PCT/US99/21090

PRIOR FILING DATE: 1999-09-15

PRIOR APPLICATION NUMBER: PCT/US99/21547

PRIOR FILING DATE: 1999-09-15

PRIOR APPLICATION NUMBER: PCT/US99/23089

PRIOR FILING DATE: 1999-10-05

PRIOR APPLICATION NUMBER: PCT/US99/28214

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; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT FILING DATE: 2001-11-20
; PRIOR APPLICATION NUMBER: US/09/907,841
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-907-841-261

Query Match          97.6%; Score 2080; DB 9; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.2e-193;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1  MAGIPGLFLLFLLCAVGQVSPYSAPWKPETWPAYRLPVVLPQSTLNLAQPDFGAEAKLE 60
Db      1  MAGIPGLFLLFLLCAVGQVSPYSAPWKPETWPAYRLPVVLPQSTLNLAQPDFGAEAKLE 60

Qy      61  VSSSCGPQCHKGTPLTYBEAKQYLSYETLIYANGSRTEQVGIYILSSGSDGAQHRDSDGS 120
Db      61  VSSSCGPQCHKGTPLTYBEAKQYLSYETLIYANGSRTEQVGIYILSSGSDGAQHRDSDGS 120

Qy      121  SGKSRKRQIYGYDVSFISFGKDFLNYPESTSVKLTGCTGLVAEKHVLTAACHHDG 180
Db      121  SGKSRKRQIYGYDVSFISFGKDFLNYPESTSVKLTGCTGLVAEKHVLTAACHHDG 180

Qy      181  KTVYVGTQKLRVGLFKPKFGKGGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240
Db      181  KTVYVGTQKLRVGLFKPKFGKGGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND 240

Qy      241  IGMDDYALLELKKPHKPKMKI GVSPPAKQLPGGRHFSGYNDPRGNLVYRFDVQKDE 300
Db      241  IGMDDYALLELKKPHKPKMKI GVSPPAKQLPGGRHFSGYNDPRGNLVYRFDVQKDE 300

Qy      301  TYDLLYQCCDAQPGASGSGVYVVMKRRQKQKWERKLIIGIFSGHQWDMNGSPQDFNVAVR 360
Db      301  TYDLLYQCCDAQPGASGSGVYVVMKRRQKQKWERKLIIGIFSGHQWDMNGSPQDFNVAVR 360
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Qy      361  ITPLYAQICYWIKGNYLDCREG 383
Db      361  ITPLYAQICYWIKGNYLDCREG 383

RESULT 8
US-09-904-011-261
; Sequence 261, Application US/09904011
; Publication No. US20030003530A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/904,011
; PRIOR FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
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;; PRIOR FILING DATE: 1999-12-20
;; PRIOR APPLICATION NUMBER: PCT/US00/00219
;; PRIOR FILING DATE: 2000-01-05
;; NUMBER OF SEQ ID NOS: 423
;; SEQ ID NO 261
;; LENGTH: 383
;; TYPE: PRT
;; ORGANISM: Homo Sapien
US-09-904-011-261

Query Match 97.6%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.2e-193;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPOCHKGTPLPTYEAKQYLSYETLYANGSRTETQVGIIYILSSSGDGAQHRDGS 120
DB 61 VSSSCGPOCHKGTPLPTYEAKQYLSYETLYANGSRTETQVGIIYILSSSGDGAQHRDGS 120
QY 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLTGCTGTVAEKHVLTAACHIDG 180
DB 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLTGCTGTVAEKHVLTAACHIDG 180
QY 181 KTVVGTQKLRVGLFKPKFQDGGGANDSTSAMPEQMKFQWIRVKTHTVPKGIKNAND 240
DB 181 KTVVGTQKLRVGLFKPKFQDGGGANDSTSAMPEQMKFQWIRVKTHTVPKGIKNAND 240
QY 241 IGMDDYDYLLELKKPKHKKFMKIGVSPPAKQLPGGRIHFSGYNDNDRPGLNLYRFCVDKDE 300
DB 241 IGMDDYDYLLELKKPKHKKFMKIGVSPPAKQLPGGRIHFSGYNDNDRPGLNLYRFCVDKDE 300
QY 301 TYDLYQQCDAOPGASGSGVYVMMKRRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
DB 301 TYDLYQQCDAOPGASGSGVYVMMKRRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPLKYAQICYWIKGNLYDCREG 383
DB 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 9
US-09-903-640-261
;; Sequence 261, Application US/09903640
;; Publication No. US20030017463A1
;; GENERAL INFORMATION:

;; APPLICANT: Genentech, Inc.
;; APPLICANT: Ashkenazi, Avi
;; APPLICANT: Botstein, David
;; APPLICANT: Desnoyers, Luc
;; APPLICANT: Eaton, Dan L.
;; APPLICANT: Ferrara, Napoleone
;; APPLICANT: Filvaroff, Ellen
;; APPLICANT: Fong, Sherman
;; APPLICANT: Gao, Wei-Qiang
;; APPLICANT: Gerber, Hanspeter
;; APPLICANT: Gerritsen, Mary E.
;; APPLICANT: Goddard, A.
;; APPLICANT: Godowski, Paul J.
;; APPLICANT: Grimaldi, Christopher J.
;; APPLICANT: Hillan, Kenneth, J.
;; APPLICANT: Kljavin, Ivar J.
;; APPLICANT: Mather, Jennie P.
;; APPLICANT: Pan, James
;; APPLICANT: Paoni, Nicholas F.
;; APPLICANT: Roy, Margaret Ann
;; APPLICANT: Stewart, Timothy A.
;; APPLICANT: Tumas, Daniel
;; APPLICANT: Williams, P. Mickey
;; APPLICANT: Wood, William, I.

;; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;; TITLE OF INVENTION: Acids Encoding the Same
;; FILE REFERENCE: 10466-14
;; CURRENT APPLICATION NUMBER: US/09/903,640
;; CURRENT FILING DATE: 2001-07-11
;; PRIOR APPLICATION NUMBER: 09/665,350
;; PRIOR FILING DATE: 2000-09-18
;; NUMBER OF SEQ ID NOS: 423
;; SEQ ID NO 261
;; LENGTH: 383
;; TYPE: PRT
;; ORGANISM: Homo Sapien
US-09-903-640-261

Query Match 97.6%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.2e-193;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPOCHKGTPLPTYEAKQYLSYETLYANGSRTETQVGIIYILSSSGDGAQHRDGS 120
DB 61 VSSSCGPOCHKGTPLPTYEAKQYLSYETLYANGSRTETQVGIIYILSSSGDGAQHRDGS 120
QY 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLTGCTGTVAEKHVLTAACHIDG 180
DB 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLTGCTGTVAEKHVLTAACHIDG 180
QY 181 KTVVGTQKLRVGLFKPKFQDGGGANDSTSAMPEQMKFQWIRVKTHTVPKGIKNAND 240
DB 181 KTVVGTQKLRVGLFKPKFQDGGGANDSTSAMPEQMKFQWIRVKTHTVPKGIKNAND 240
QY 241 IGMDDYDYLLELKKPKHKKFMKIGVSPPAKQLPGGRIHFSGYNDNDRPGLNLYRFCVDKDE 300
DB 241 IGMDDYDYLLELKKPKHKKFMKIGVSPPAKQLPGGRIHFSGYNDNDRPGLNLYRFCVDKDE 300
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DB 301 TYDLYQQCDAOPGASGSGVYVMMKRRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPLKYAQICYWIKGNLYDCREG 383
DB 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 10
US-09-908-093-261
;; Sequence 261, Application US/09908093
;; Publication No. US20030017498A1
;; GENERAL INFORMATION:

;; APPLICANT: Genentech, Inc.
;; APPLICANT: Ashkenazi, Avi
;; APPLICANT: Botstein, David
;; APPLICANT: Desnoyers, Luc
;; APPLICANT: Eaton, Dan L.
;; APPLICANT: Ferrara, Napoleone
;; APPLICANT: Filvaroff, Ellen
;; APPLICANT: Fong, Sherman
;; APPLICANT: Gao, Wei-Qiang
;; APPLICANT: Gerber, Hanspeter
;; APPLICANT: Gerritsen, Mary E.
;; APPLICANT: Goddard, A.
;; APPLICANT: Godowski, Paul J.
;; APPLICANT: Grimaldi, Christopher J.
;; APPLICANT: Gurney, Austin L.
;; APPLICANT: Hillan, Kenneth, J.
;; APPLICANT: Kljavin, Ivar J.
;; APPLICANT: Mather, Jennie P.
;; APPLICANT: Pan, James
;; APPLICANT: Paoni, Nicholas F.
;; APPLICANT: Roy, Margaret Ann

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; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/908,093
; CURRENT FILING DATE: 2001-07-17
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-908-093-261

Query Match          97.6%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 2,2e-193;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1  MAGIPGLLFLLFLLCAVGQSPYSAPWKPPTWPAYRLPVVLPOSTLNLAKEPFGAEAKLE 60
      |||
Db      1  MAGIPGLLFLLFLLCAVGQSPYSAPWKPPTWPAYRLPVVLPOSTLNLAKEPFGAEAKLE 60
      |||

QY      61  VSSSCGPOCHKGTPPTYPEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDGS 120
      |||
Db      61  VSSSCGPOCHKGTPPTYPEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDGS 120
      |||

QY      121  SGKSRKROIYGVDSRFSIFGKDFLNYPSTSVKLSCTGCTFLVAEKHVLTAACHIDG 180
      |||
Db      121  SGKSRKROIYGVDSRFSIFGKDFLNYPSTSVKLSCTGCTFLVAEKHVLTAACHIDG 180
      |||

QY      181  KTVVKGTQKLVRGFLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
      |||
Db      181  KTVVKGTQKLVRGFLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
      |||

QY      241  IGMDDYALLELEKKPHKRFMKIGVSPPAKQLPGGRIHFSGYDNDRPGNLVYRFCVDKDE 300
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Db      241  IGMDDYALLELEKKPHKRFMKIGVSPPAKQLPGGRIHFSGYDNDRPGNLVYRFCVDKDE 300
      |||
QY      301  TYDLLYQQCDAQGASGSGVYVWMKRCQQKWKRIIGIFSGHQWYDMNGSPQDFNVAVR 360
      |||
Db      301  TYDLLYQQCDAQGASGSGVYVWMKRCQQKWKRIIGIFSGHQWYDMNGSPQDFNVAVR 360
      |||
QY      361  ITPKYAQICYWIKGNYLDCREG 383
      |||
Db      361  ITPKYAQICYWIKGNYLDCREG 383
      |||

RESULT 11
US-09-906-742-261
; Sequence 261, Application US/09906742
; Publication No. US20030023054A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/906,742
; CURRENT FILING DATE: 2001-07-16
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
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; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-906-742-261

Query Match          97.6%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.2e-193;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLA KPDPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLA KPDPFGAEAKLE 60
QY 61 VSSSCGPGCHKGTPLTPEEAKQYLSYETLYANGSRSTETQVGIIYILSSSGDGAQHRS DGS 120
Db 61 VSSSCGPGCHKGTPLTPEEAKQYLSYETLYANGSRSTETQVGIIYILSSSGDGAQHRS DGS 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGTLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGTLVAEKHVLTAACHIDG 180
QY 181 KTYVGTQKLRVGLFKPKFGKGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
Db 181 KTYVGTQKLRVGLFKPKFGKGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
QY 241 IGHVDYALLELKKPKHKFKMKIGVSPPAKQLPGGRHSHSGYDNDPRGNLVYRFCVDKDE 300
Db 241 IGHVDYALLELKKPKHKFKMKIGVSPPAKQLPGGRHSHSGYDNDPRGNLVYRFCVDKDE 300
QY 301 TYDLLYQQCDAQPGASGSGVYVVMKRWQKQKWKRIIGIFSGHGWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAQPGASGSGVYVVMKRWQKQKWKRIIGIFSGHGWDMNGSPQDFNVAVR 360
QY 361 ITPLKVAQICYWIKGNLYDCREG 383
Db 361 ITPLKVAQICYWIKGNLYDCREG 383

RESULT 12
US-09-906-838-261
; Sequence 261, Application US/09906838
; Publication No. US20030027143A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnovers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanepeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavini, Ivar J.
; APPLICANT: Mather, Jennie P.
```

```
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas P.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/906,838
; CURRENT FILING DATE: 2001-07-16
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
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; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-906-838-261

Query Match          97.6%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.2e-193;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLA KPDPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLA KPDPFGAEAKLE 60
QY 61 VSSSCGPGCHKGTPLTPEEAKQYLSYETLYANGSRSTETQVGIIYILSSSGDGAQHRS DGS 120
Db 61 VSSSCGPGCHKGTPLTPEEAKQYLSYETLYANGSRSTETQVGIIYILSSSGDGAQHRS DGS 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGTLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGTLVAEKHVLTAACHIDG 180
QY 181 KTYVGTQKLRVGLFKPKFGKGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
Db 181 KTYVGTQKLRVGLFKPKFGKGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
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Db 181 KTYVGTQKLRVGLFKPKFKDGGGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
QY 241 IGMDDYDYLLELKKPKFKFKMKGIVSPPAKQLFGGRHFSGYDNDPRGNLVYRFCDDVDE 300
Db 241 IGMDDYDYLLELKKPKFKFKMKGIVSPPAKQLFGGRHFSGYDNDPRGNLVYRFCDDVDE 300
QY 301 TYDLLYQQCDAOPGASGSGVYVMMKROQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAOPGASGSGVYVMMKROQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPKYAQICYWIKNYLDREG 383
Db 361 ITPKYAQICYWIKNYLDREG 383

RESULT 13
US-09-907-613-261
; Sequence 261, Application US/09907613
; Publication No. US20030027145A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,613
; PRIOR FILING DATE: 2001-07-17
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564

; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; TYPE: PRT
; LENGTH: 383
; ORGANISM: Homo Sapien
US-09-907-613-261
Query Match 97.6%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.2e-193;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MAGIPQLLFLFLCAVGVSPYSAPMKPTWPAYRLPVVLQSTLNLAKPFGAEAKLE 60
Db 1 MAGIPQLLFLFLCAVGVSPYSAPMKPTWPAYRLPVVLQSTLNLAKPFGAEAKLE 60
QY 61 VSSSCGPQCHKGTPLTPTVEEAKQYLSYETLYANGSRTETQVGIIYILSSSGDGAQHRDGS 120
Db 61 VSSSCGPQCHKGTPLTPTVEEAKQYLSYETLYANGSRTETQVGIIYILSSSGDGAQHRDGS 120
QY 121 SKSRRKRIQYDYSRFSIFGKDFLLNYPFSTSVKLSCTCTGLVAEKHVLTAACHIDG 180
Db 121 SKSRRKRIQYDYSRFSIFGKDFLLNYPFSTSVKLSCTCTGLVAEKHVLTAACHIDG 180
QY 181 KTYVGTQKLRVGLFKPKFKDGGGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
Db 181 KTYVGTQKLRVGLFKPKFKDGGGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
QY 241 IGMDDYDYLLELKKPKFKFKMKGIVSPPAKQLFGGRHFSGYDNDPRGNLVYRFCDDVDE 300
Db 241 IGMDDYDYLLELKKPKFKFKMKGIVSPPAKQLFGGRHFSGYDNDPRGNLVYRFCDDVDE 300
QY 301 TYDLLYQQCDAOPGASGSGVYVMMKROQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAOPGASGSGVYVMMKROQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPKYAQICYWIKNYLDREG 383
Db 361 ITPKYAQICYWIKNYLDREG 383
RESULT 14
US-09-907-942-261
; Sequence 261, Application US/09907942
; Publication No. US20030027146A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavin, Ivar J.

APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/907,942
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR FILING DATE: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/30095
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-907-942-261

Query Match 97.6%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 2,2e-193;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCVAGQVSPYAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLLCVAGQVSPYAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGQCHKGTPLTPTVEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS 120
DB 61 VSSSCGQCHKGTPLTPTVEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS 120
QY 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSGTCTGLTVAEKVLTAAHCHD 180
DB 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSGTCTGLTVAEKVLTAAHCHD 180
QY 181 KTVVKGTKLRVGLFKPKDGGGRGNDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
DB 181 KTVVKGTKLRVGLFKPKDGGGRGNDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240

QY 241 IGMVDYDYLLELKKPHKRFMKIGVSPPAKQLPGGRIHESGYDNDPRGNLVYRFDVKDE 300
DB 241 IGMVDYDYLLELKKPHKRFMKIGVSPPAKQLPGGRIHESGYDNDPRGNLVYRFDVKDE 300
QY 301 TYDLLLYQQCDAQPGASGSGVYVYRMWKRQQQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
DB 301 TYDLLLYQQCDAQPGASGSGVYVYRMWKRQQQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPLKYAQICYWIKGNLYDCREG 383
DB 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 15
US-09-904-859-261
Sequence 261, Application US/09904859
Publication No. US20030036060A1
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/904,859
CURRENT FILING DATE: 2001-07-12
PRIOR APPLICATION NUMBER: 09/665,350
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30

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; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-904-859-261

Query Match          97.6%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 2.2e-193;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1  MAGIPGLLFLFLLCAVGVSPYSAPWKPWTWPAYRLPVVLPQSTLNLAKPDPFGAEAKLE 60
Db      1  MAGIPGLLFLFLLCAVGVSPYSAPWKPWTWPAYRLPVVLPQSTLNLAKPDPFGAEAKLE 60

Qy      61  VSSSCGPGCHKGTPPTYEAKQYLSYETLYANGSRTEQVGIYIILSSGDDGAQHRDGS 120
Db      61  VSSSCGPGCHKGTPPTYEAKQYLSYETLYANGSRTEQVGIYIILSSGDDGAQHRDGS 120

Qy      121  SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSTCTGTVAEKHVLTAACHIDG 180
Db      121  SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSTCTGTVAEKHVLTAACHIDG 180

Qy      181  KTVVKTOKLRVGFLLKPKFKDGGGANDSTSAMPEQMKQWIRVKETHVPKGWIKGNAND 240
Db      181  KTVVKTOKLRVGFLLKPKFKDGGGANDSTSAMPEQMKQWIRVKETHVPKGWIKGNAND 240

Qy      241  IGMDYDVALLELKKPKHKRFMKIGVSPPAKQLPGGRIHFSGYDNDPRGNLVYRFCVDKDE 300
Db      241  IGMDYDVALLELKKPKHKRFMKIGVSPPAKQLPGGRIHFSGYDNDPRGNLVYRFCVDKDE 300

Qy      301  TYDLLYQQCDAQPGASGSGYVVMWKRQQQKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Db      301  TYDLLYQQCDAQPGASGSGYVVMWKRQQQKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360

Qy      361  ITPKYAQICYWIKGNLYDCREG 383
Db      361  ITPKYAQICYWIKGNLYDCREG 383
```

Search completed: July 1, 2005, 21:32:17
Job time : 91.3582 secs

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: July 1, 2005, 20:54:23 ; Search time 20.826 Seconds
(without alignments)
1811.048 Million cell updates/sec

Title: US-09-658-677-15
Perfect score: 2131
Sequence: 1 MAGIPGLLFLFLCAVGO.....IKGNLYDCREGDTVPFGSN 392

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues
Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : PIR 79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	194	9.1	316	2 A45134	endopeptidase (EC
2	135.5	6.4	313	2 A35122	metalloproteinase
3	131	6.1	218	2 E97915	choline binding pr
4	120.5	5.7	482	1 EXRT	coagulation factor
5	119	5.6	269	2 A26823	pancreatic elastas
6	117	5.5	522	2 T29767	hypothetical prote
7	115.5	5.4	269	2 B26823	pancreatic elastas
8	115.5	5.4	271	2 A25528	pancreatic elastas
9	115	5.4	258	4 S70439	pancreatic elastas
10	115	5.4	267	4 A56615	probable pancreati
11	114.5	5.4	238	1 TRW5Y	trypsin-like prote
12	114.5	5.4	285	2 C95045	choline binding pr
13	114	5.3	246	1 DBHU	complement factor
14	113	5.3	266	1 ELPG	pancreatic elastas
15	113	5.3	266	1 ELRT1	pancreatic elastas
16	112.5	5.3	259	1 TRSMG	trypsin (EC 3.4.21
17	112.5	5.3	273	2 E85765	hypothetical prote
18	112.5	5.3	273	2 H64915	putative protease
19	111.5	5.2	488	1 EXHU	coagulation factor
20	111	5.2	761	2 JC5759	brain-specific ser
21	110	5.2	269	2 C26823	pancreatic elastas
22	110	5.2	492	1 EXBO	coagulation factor
23	110	5.2	1582	2 T15308	hypothetical prote
24	109.5	5.1	405	2 T35117	probable secreted
25	109	5.1	278	2 A20282	probable peptidas
26	108	5.1	583	2 A29154	complement factor
27	108	5.1	786	1 A47547	serine proteinase
28	108	5.1	1047	2 A55617	masquerade precurs
29	107.5	5.0	236	2 A28566	T-cell suppressor

30	107.5	5.0	686	1 A59271	Ra-reactive factor
31	105	4.9	274	2 S40004	trypsin-related pr
32	103.5	4.9	271	1 ELRT2	pancreatic elastas
33	103.5	4.9	416	1 KFB0	coagulation factor
34	102.5	4.8	1238	2 T34929	hypothetical prote
35	100	4.7	272	2 JC4170	trypsin-like prote
36	100	4.7	855	2 JC7731	membrane-bound arg
37	98.5	4.6	267	2 S40006	trypsin (EC 3.4.21
38	98	4.6	548	2 D82175	probable trypsin V
39	98	4.6	624	2 T02289	probable polygalac
40	97.5	4.6	409	2 T35118	probable secreted
41	97.5	4.6	452	1 A30351	coagulation factor
42	97.5	4.6	747	2 I51579	complement factor
43	97	4.6	1019	2 A38738	coagulation factor
44	96.5	4.5	259	2 S68424	allergen Der f III
45	96.5	4.5	275	2 I46712	factor IX - rabbit

ALIGNMENTS

RESULT 1

A45134
endopeptidase (EC 3.4.-.-), glutamate-specific - Bacillus licheniformis
C;Species: Bacillus licheniformis
C;Date: 10-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 16-Aug-2004
C;Accession: A45134; S23078
R;Kakudo, S.; Kikuchi, N.; Kitadokoro, K.; Fujiwara, T.; Nakamura, E.; Okamoto, H.; Shin, J. Biol. Chem. 267, 23782-23788, 1992
A;Title: Purification, characterization, cloning, and expression of a glutamic acid-speci
A;Reference number: A45134; MUID:93054737; PMID:1429718
A;Accession: A45134
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-316 <RAK>
A;Cross-references: UNIPROT:P80057; GB:D10060; NID:g216263; PIDN:BAA00949.1; PID:d100141;
A;Experimental source: ATCC 14580
A;Note: sequence extracted from NCBI backbone (NCBIN:118784, NCBIPI:118785)
R;Svendsen, I.; Breddam, K.
Eur. J. Biochem. 204, 165-171, 1992
A;Title: Isolation and amino acid sequence of a glutamic acid specific endopeptidase from
A;Reference number: S23078; MUID:92155199; PMID:1346764
A;Accession: S23078
A;Status: preliminary
A;Molecule type: protein
A;Residues: 95-316 <SVB>
C;Superfamily: Glutamyl endopeptidase, V8 type
C;Keywords: hydrolase

Query Match 9.1%; Score 194; DB 2; Length 316;
Best Local Similarity 24.0%; Pred. No. 2e-08;
Matches 79; Conservative 43; Mismatches 143; Indels 64; Gaps 15;

QY	63	SSGPGCHGTPL---PTVEEAKQVLSYETLVANGSRTEQTQVGIYILSSSGDGAQRHDSG	119
DB	28	AQAPSPH--TPVSSDPST-KAETSVDYD-----NIKSDQYGLYSKAFGTGKVNETKE	79
QY	120	SSGSKRRKQIYGYDSRFRIFGKDFLLN-----YPFSTSVKLST---GCTGLVAEKHV	170
DB	80	KAEKSPAKAPY---SIKSVIGSDDRTRVTNTTAYPYRAIVHISSTSGCTGWMIGPKTV	136
QY	171	LTNAHCIHGKGT-YVKGTKLRVGLFKPKFKDGGGRANDSTSAMPEQMKQWIRVKRTHV	229
DB	137	ATAGHCIIYDTSSGSPAGTATVSPG-----RNGTS-----YYPGSKVSTRFYI	178
QY	230	PKGWIKGNANDIGMDYDVALLELKPDKRKFMIKIGVSPPAKQLPGGRIHFSGYDNDNRPGN	289
DB	179	PSGWRSGNTN-----YDYGAIELSEFIQNTVGVFGYSYTTSSLVGTTVTISGPGDKTAG	233
QY	290	LVYRFCD--VKDETLDLYQQCDAPQASGSGVYVVRMKRQQQKWERKII---GIFSGH	343
DB	234	TQWHSPTAISETYKLYQYAM-DTYGQSGSPVFEQSSSRSTNCSGPCSLAVHTNGVYGG-	291

```
QY 344 QWDMNGSPQDFNVAVRITPLKVAQICYW 372
Db 292 -----SSYNRGTRITKEVFDNLTNW 311

RESULT 2
A35122
metalloprotease (EC 3.4.21.6) mpr precursor, extracellular - Bacillus subtilis
C:Species: Bacillus subtilis
C>Date: 27-Jul-1990 #sequence_revision 27-Jul-1990 #text_change 16-Aug-2004
C:Accession: A35122; I40010; A69660
R:Sloma, A.; Rudolph, C.F.; Rufo Jr., G.A.; Sullivan, B.J.; Theriault, K.A.; Ally, D.; F
J. Bacteriol. 172, 1024-1029, 1990
A:Title: Gene encoding a novel extracellular metalloprotease in Bacillus subtilis.
A:Reference number: A35122; MUID:90130256; PMID:2105291
A:Status: preliminary
A:Accession: A35122
A:Molecule type: DNA
A:Residues: 1-313 <SLO>
A:CROSS-references: UNIPROT:P39790; GB:L10505; NID:g143209; PIDN:AAA22604.1; PID:g143210
R:Smith, H.; de Jong, A.; Bron, S.; Venema, G.
Gene 70, 351-361, 1988
A:Title: Characterization of signal-sequence-coding regions selected from the Bacillus s
A:Reference number: I39994; MUID:89108019; PMID:3145906
A:Accession: I40010
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-60, 65, 'L', 67, 'S', 69, 'AQA' <RES>
A:CROSS-references: GB:M22916; NID:g143701; PIDN:AAA22832.1; PID:g143702
R:Kunst, F.; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Bertel
C.; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Ch
A.; Ehrlich, S.D.; Emmeron, P.T.; Entian, K.D.; Errington, J.; Fabbret, C.; Ferrari, E.
Nature 390, 249-256, 1997
A:Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Gallen
iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holzapfel, S.; Hosono, S.; Hullo, M.F.
Koetter, P.; Koningsstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois,
A:Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maueel
Y, M.; Ogawa, K.; Ogilwara, A.; Oudeg, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetelle
Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon,
A:Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Seron
akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchiyama
T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, K
A:Authors: Yoshikawa, H.F.; Zumbstein, E.; Yoshikawa, H.; Danchin, A.
A:Title: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis.
A:Reference number: A69580; MUID:98044033; PMID:9384377
A:Accession: A69660
A:Status: preliminary; nucleic acid sequence not shown; translation not shown
A:Molecule type: DNA
A:Residues: 1-313 <KUN>
A:CROSS-references: GB:Z99105; GB:AL009126; NID:g2632457; PIDN:CAB12018.1; PID:el182176;
A:Experimental source: strain 168
C:Genetics:
A:Gene: mpr
C:Superfamily: Glutamyl endopeptidase, V8 type
C:Keywords: hydrolase

Query Match 6.4%; Score 135.5; DB 2; Length 313;
Best Local Similarity 22.2%; Pred. No. 0.0014;
Matches 77; Conservative 38; Mismatches 121; Indels 111; Gaps 17;

QY 72 GTPLPYEEAKQVLSVETLYANGSRTRTQVGIYILSSGD--CAQHRDSSGSKSRKQ 129
Db 29 GVPKAAENPQTSVNTSGREATKNQT-----SKADQVSAPYEGTGKTSK----- 75
QY 130 IXYGDSRF-----SIFGKD-----FLANYPFSTSVKLST-----GCTGTL 164
Db 76 LYGGQTELEKNIGTLOFSSIIIGTDERTSSTTSFPYRATVQLSIKYPTNSYTGCTGFL 135
QY 165 VAEKHVLTAAHCIIH-----DGKTYVKGQKLRVGLFKPKFKDGGRGAND 208
Db 136 VNPNVTVTAGHCYVSQDHGWASTITAAPGRNGSSYPYGIY----- 175
QY 209 STSAMPEQMKFQIRVKRTHRVPGWIKGN---ANDIGMDYDVALLELKKPKFKFKMKIGV 265
```

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Db 176 -SCTMFYSVK-CWTESKDTNYDYGAIKLNGSPGNTVGM-YGYRTTNSSGP-----VGL 225
QY 266 SPPAKQLPGGRIFHSYGDNDPRGNLVYRFDCKVDETYDLLYQOCDAQPGASGVVVRMW 325
Db 226 SSSVVGFPDCKTFGTWMSDTKPIR-----SAETYKLTYY-TTDTYGCQSGSPVY---- 272
QY 326 KRQQQKWKERKIIGIFSGHQMVDNMGSPQDFNVAVRITPLKVAQICYW 372
Db 273 -RNYSDTGQTALAIHT-----NGG-SSYNLGTRVNDVFNNIQYW 310

RESULT 3
E97915
choline binding protein G, truncation [imported] - Streptococcus pneumoniae (strain R6)
C:Species: Streptococcus pneumoniae
C>Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 09-Jul-2004
C:Accession: E97915
R:Hoskins, J.A.; Alborn Jr., W.; Arnold, J.; Blaszcak, L.; Burgett, S.; DeHoff, B.S.; E
e, R.; LeBlanc, D.J.; Lee, L.N.; Lefkowitz, E.J.; Lu, J.; Matsushima, P.; McAhren, S.; M
Y, P.; Sun, P.M.; Winkler, M.E.
J. Bacteriol. 183, 5709-5717, 2001
A:Authors: Yang, Y.; Young-Bellido, M.; Zhao, G.; Zook, C.; Baltz, R.H.; Jaskunas, S.R.;
A:Title: Genome of the Bacterium Streptococcus pneumoniae Strain R6.
A:Reference number: A97872; MUID:21429245; PMID:11544234
A:Accession: E97915
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-218 <KUR>
A:CROSS-references: UNIPROT:Q8DR41; GB:AE007317; PIDN:AAK99153.1; PID:g15457907; GSPDB:GB
C:Genetics:
A:Gene: cbpG-truncation

Query Match 6.1%; Score 131; DB 2; Length 218;
Best Local Similarity 24.2%; Pred. No. 0.0021;
Matches 59; Conservative 30; Mismatches 87; Indels 68; Gaps 10;

QY 143 DFLNYPFSTSVKLSGTGCTG-----TLVAEKHVLTAACHIDHGTYYKGTOK 189
Db 5 DNTLQVPYSTSAWLSKYVGVADGMNVEGRSGSANFKDNLVITAAH-----NYR---- 54
QY 190 LRVGFLKPKFKDGGRGAND---STSAMPEQMKFQIRVKRTHVPGWIKGNANDIGMDYD 246
Db 55 -----HDYGEADIIYVLPVAVSPSQELFKYKVEVRYLKEFRNLNSKD-AREYD 103
QY 247 YALLELKKPKFKFKMGKIVGSPPAKOLPGGRIFHSYGDNDPRGNLVYR-----CDVKDE 300
Db 104 LALLILEKPIGAKLTGLTGLTSGKSLTGITVITGYPS-----YNFKIHQMYTDKKQV 156
QY 301 TYD---LLYQOCDAQPGASGVVYRMMKROQKWKERKIIGIFSGHQMVDNMGSPQDFNV 357
Db 157 LSDDGMFLDYQVDTLEGSSGSTVY-----DASHRVVGVHT-----LGDGANQINS 201
QY 358 AVRI 361
Db 202 AVKL 205

RESULT 4
EXPT
coagulation factor Xa (EC 3.4.21.6) precursor - rat
C:Species: Rattus norvegicus (Norway rat)
C>Date: 31-Jan-1995 #sequence_revision 07-Feb-1997 #text_change 09-Jul-2004
C:Accession: S49075; JC4670; PS0191; PS0190; I62745
R:Stanton, C.; Ross, P.; Hutson, S.; Wallin, R.
Thromb. Res. 80, 63-73, 1995
A:Title: Evidence for competition between vitamin K-dependent clotting factors for intrac
A:Reference number: A59498; MUID:96093366; PMID:8578539
A:Accession: S49075
A:Molecule type: mRNA
A:Residues: 1-482 <STAL>
A:CROSS-references: UNIPROT:Q63207; EMBL:X79807; NID:g506600; PIDN:CAA56202.1; PID:g50660
A:Note: submitted to the EMBL Data Library, June 1994
```

A>Note: neither the complete nucleic acid sequence nor the complete translation are shown
R:Stanton, C.; Ross, R.P.; Hutson, S.; Wallin, R.
Gene 159, 269-273, 1996
A>Title: Processing and expression of rat and human clotting factor-X-encoding cDNAs.
A:Reference number: JC4670; MUID:96194815; PMID:8647460
A:Molecule type: mRNA
A:Accession: JC4670
A:Residues: 1-482 <STA2>
A:Cross-references: EMBL:X79807; NID:G506600; PIDN:CAA56202.1; PID:G506601
A:Experimental source: Cos-1 cell
R:Enjoji, K.; Miyazaki, K.; Kato, H.
J. Biochem. 109, 890-898, 1991
A>Title: Characterization of rat factors X and Xa: demonstration of factor Xa in rat plasma
A:Reference number: PS0190; MUID:92041742; PMID:1718949
A:Accession: PS0191
A:Molecule type: protein
A:Residues: 41-58, 'X', 60-65 <ENJ1>
A:Accession: PS0190
A:Molecule type: protein
A:Residues: 183-186, 'X', 188-207 <ENJ2>
R:Murakawa, M.; Okamura, T.; Kamura, T.; Kuroiwa, M.; Harada, M.; Niho, Y.
Eur. J. Haematol. 52, 162-168, 1994
A>Title: Analysis of the partial nucleotide sequences and deduced primary structures of factor Xa
A:Reference number: I46196; MUID:94222160; PMID:8168596
A:Accession: I62745
A:Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: DNA
A:Residues: 295-383, 'G', 385-455 <MUR>
A:Cross-references: GB:D21215; NID:G415309; PIDN:BAA04756.1; PID:G455396
C:Function:
A:Description: catalyzes the proteolytic activation of prothrombin to thrombin in the presence of calcium ions
A:Pathway: blood coagulation
C:Superfamily: coagulation factor X; EGF homology; Gla domain homology; trypsin homology
C:Keywords: beta-hydroxyaspartic acid; blood coagulation; calcium binding; carboxyglutamate
F:1-23/Domain: signal sequence #status predicted <SIG>
F:24-40/Domain: propeptide #status predicted <PRO>
F:25-84/Domain: Gla domain homology <GLA>
F:41-179/Product: coagulation factor X light chain #status predicted <LCH>
F:90-121/Domain: EGF homology <EGF>
F:129-164/Domain: EGF homology <EG2>
F:183-482/Product: coagulation factor X heavy chain #status predicted <HCH>
F:183-231/Domain: activation peptide #status predicted <APT>
F:232-482/Product: coagulation factor Xa heavy chain #status predicted <ACT>
F:232-460/Domain: trypsin homology <TRY>
F:46,47,54,56,59,60,65,66,69,72,79/Modified site: gamma-carboxyglutamic acid (Glu) #status predicted <MOD>
F:57-62,90-101,95-110,112-121,129-140,136-149,151-164,172-340,238-243,259-275,388-402,411-421/Modified site: erythro-beta-hydroxyaspartic acid (Asp) #status predicted <MOD>
F:103/Modified site: carboxy-beta-hydroxyaspartic acid (Asp) #status predicted <MOD>
F:187/Binding site: carboxy-beta (Asn) (covalent) #status experimental
F:208/Binding site: carboxy-beta (Thr) (covalent) #status predicted
F:218/Binding site: carboxy-beta (Asn) (covalent) #status predicted
F:231-232/Cleavage site: Arg-Ile (coagulation factor IXa, coagulation factor VIIa) #status predicted
F:274,320,417/Active site: His, Asp, Ser #status predicted

Query Match 5.7%; Score 120.5; DB 1; Length 482;
Best Local Similarity 32.3%; Pred. No. 0.041;
Matches 31; Conservative 17; Mismatches 23; Indels 25; Gaps 4;
A:Introns: 36/2; 138/3; 234/2; 311/1; 331/1; 421/1; 470/2

QY 160 CTGTLVAEKHLTAACHIDHDKTYVKGTKLVGFLKPKFGDGRGANDSTSAMPEOMKF 219
DB 259 CGGTILNEFYLTAAHCLHQAQRF-----KVRVGDNLTEQEDGGMVHE-VDMLIKHNKF 312

QY 220 QWTRVVRKTHVPKWKGNANDIGMDYDYLLEKKP 255
DB 313 Q-----RDTY-----DFDIAMLRKTP 329

RESULT 5
A26823
pancreatic elastase II (EC 3.4.21.71) precursor - pig
N:Alternate names: pancratopeptidase E
C:Species: Sus scrofa domestica (domestic pig)
C:Date: 16-Aug-1988 #sequence_revision 16-Aug-1988 #text_change 09-Jul-2004
C:Accession: A26823

R:Kawashima, I.; Tani, T.; Shimoda, K.; Takiguchi, Y.
DNA 6, 163-172, 1987
A>Title: Characterization of pancreatic elastase II cDNAs: two elastase II mRNAs are expressed in the pancreas of the rat
A:Reference number: A90958; MUID:87217962; PMID:3646943
A:Accession: A26823
A:Molecule type: mRNA
A:Residues: 1-269 <KAW>
A:Cross-references: UNIPROT:P08419; GB:M16651; NID:G164441; PIDN:AAA31027.1; PID:G164442
C:Superfamily: trypsin; trypsin homology
C:Keywords: hydrolase; serine proteinase
F:1-16/Domain: signal sequence #status predicted <SIG>
F:17-28/Domain: propeptide #status predicted <PRO>
F:29-269/Product: elastase II #status predicted <MAT>
F:29-262/Domain: trypsin homology <TRY>
F:73,121,216/Active site: His, Asp, Ser #status predicted

Query Match 5.6%; Score 119; DB 2; Length 269;
Best Local Similarity 27.5%; Pred. No. 0.027;
Matches 60; Conservative 28; Mismatches 64; Indels 66; Gaps 14;
A:Introns: 36/2; 138/3; 234/2; 311/1; 331/1; 421/1; 470/2

QY 128 RQIYGYDYSFSGKDFLLNYPSTSVKL-STG-----CTGTLVAEKHLTAACHIDGK 181
DB 28 RVVGGEDARPN-----SWPMQVSLQYDSSQWRHTCGTGLVDQSWLTAACHICSSR 79

QY 182 TY--VKGTKLVGFLKPKFGDGRGANDSTSAMPEOMKFQWIRVVRKTHVPKWKGNAN 239
DB 80 TRAVLGRHSL-----STNPFGLA-----VVKSLVVDQW-----NSN 114

QY 240 DIGMDYDYLLEKKP-HKRKFMKIGVSPPAK-LFG-----GRIHFGYDND--R 286
DB 115 QLSNGNDIALKLKASPVSLTDKIQGLCPAAGTILPNVYVYVGTGRLQTNGASPDILQ 174

QY 287 PGNLVTRFCVDKDETVDLLYQQCDAPGASGSGVYVRM 324
DB 175 QGOLL-----VVD-----YATC-SKPGWGWSTVKTNM 200

RESULT 6
T29767
Hypothetical protein ZC581.6 - Caenorhabditis elegans
C:Species: Caenorhabditis elegans
C:Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
C:Accession: T29767
R:Waterston, B.; Gattung, S.; Le, T.T.
Submitted to the EMBL Data Library, May 1997
A:Description: The sequence of C. elegans cosmid ZC581.
A:Reference number: Z20682
A:Accession: T29767
A:Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: DNA
A:Residues: 1-522 <WAT>
A:Cross-references: UNIPROT:O01771; EMBL:AF003134; PIDN:AAB54144.1; GSPDB:GN00019; CESP:ZC581
A:Experimental source: strain Bristol N2; clone ZC581
C:Genetics:
A:Gene: CESP:ZC581.6
A:Map position: 1
A:Introns: 36/2; 138/3; 234/2; 311/1; 331/1; 421/1; 470/2

Query Match 5.5%; Score 117; DB 2; Length 522;
Best Local Similarity 20.4%; Pred. No. 0.089;
Matches 95; Conservative 56; Mismatches 153; Indels 162; Gaps 21;
A:Introns: 36/2; 138/3; 234/2; 311/1; 331/1; 421/1; 470/2

QY 29 KPTWPAYRPLVVLVQSTLNLAQPDFAEAKLVSSS---CGPOCHKGTPLPTVEAKQ-- 83
DB 26 KPSNKAASSAPSLRKKSSSNPNKGTARSVKSPKSAIPASPTVQKEVPPVEIEKKKEE 85

QY 84 -----YLSYETLYANGSRTEQVGIYILSSGDGAHQHSDSGSGSKRRKRIYVDSRF 137
DB 86 PENQKKEAEKKL-----DRTQDDGKEYKEAEGALGVVVIKEDKAPAKMDGDYEDFGGCEF 141

QY 138 SIFGKD-----FLNYPFFSTV-----KLSTGCTGTILVAEK 168
DB 142 PFLKILEMLVWLRIFFPSKAVYNGRDASQSEAPWSVFTLYSKDSQSATTCTGTIVSPR 201

A:Description: Isolation, sequencing and characterization of 2 cDNA clones coding for tr

RESULT 13

DBHU
Complement factor D (EC 3.4.21.46) precursor [validated] - human (fragment)
N;Alternate names: adipsin; C3 convertase activator
C;Species: Homo sapiens (man)
C;Date: 28-Aug-1985 #sequence_revision 31-Dec-1992 #text_change 09-Jul-2004
C;Accession: A40197; A00936; A60571; S66645
R;White, R.T.; Damm, D.; Hancock, N.; Rosen, B.S.; Lowell, B.B.; Usher, P.; Plier, J.S.; J. Biol. Chem. 267, 9210-9213, 1992
A;Title: Human adipsin is identical to complement factor D and is expressed at high level
A;Reference number: A40197; MUID:92250520; PMID:1374388
A;Accession: A40197
A;Molecule type: mRNA
A;Residues: 1-246 <WHI>
A;Cross-references: UNIPROT:P00746; GB:M84526
R;Niemann, M.A.; Bhowm, A.S.; Bennett, J.C.; Volanakis, J.E.
Biochemistry 23, 2482-2486, 1984
A;Title: Amino acid sequence of human D of the alternative complement pathway.
A;Reference number: A00936; MUID:85000441; PMID:6383466
A;Accession: A00936
A;Molecule type: protein
A;Residues: 19-44,'G',46-51,'Q',53-75,'TH',78,'P',80-83,'XXXITIE',90-172,86-91,185-235,'
A;Note: a few residues were assigned from the previously published sequence of Reid et al
R;Miyata, T.; Oda, O.; Inagi, R.; Sugiyama, S.; Miyama, A.; Maeda, K.; Nakashima, I.; Ya
Mol. Immunol. 27, 637-644, 1990
A;Title: Molecular and functional identification and purification of complement component
A;Reference number: A60571; MUID:90370044; PMID:2395435
A;Accession: A60571
A;Molecule type: protein
A;Residues: 19-20,'XX',23-27,'XX',30-31,'XX',34,'X',36-40 <MIY>
R;Balke, N.; Holzkamp, U.; Hoerl, W.H.; Tschesche, H.
FEBS Lett. 371, 300-302, 1995
A;Title: Inhibition of degradation of human polymorphonuclear leukocytes by complement
A;Reference number: S66645; MUID:96013156; PMID:7556615
A;Accession: S66645
A;Status: preliminary
A;Molecule type: protein
A;Residues: 19-44,'C',46-48 <BAL>
C;Comment: Factor D cleaves factor B when the latter is complexed with factor C3b, activ
A;Genetics:
A;Gene: GDB:DF
A;Cross-references: GDB:132645; OMIM:134350
A;Map position: Xpter-Xqter
C;Superfamily: trypsin; trypsin homology
C;Keywords: complement alternate pathway; hydrolase; plasma; serine proteinase
F;1-18/Domain: signal sequence #status predicted <SIG>
F;19-246/Product: complement factor D (fragment) #status experimental <MAT>
F;19-241/Domain: trypsin homology <TRY>
F;44-60,141-207,172-188,197-222/bisulfide bonds: #status predicted
F;59,105,201/Active site: His, Asp, Ser #status predicted
Query Match 5.3%; Score 114; DB 1; Length 246;
Best Local Similarity 28.4%; Pred. No. 0.062;
Matches 44; Conservative 20; Mismatches 43; Indels 48; Gaps 9;
QY 149 PFSTSVKLSGTG--CTGTLVAEKHVLTAACIHH---DGKTVYKGTQKLRVGFLLPKPKDGG 203
Db 31 PYMASVOLNGAHLCAAGVLAERWLSAAHCLDAADGKQVQL----- 72
QY 204 RGNADSTSAMPEQKQWIRVKTTHVPKGIKGNANDIGMDYDVALLELKPKHKPKMKI 263
Db 73 LGAHSLSQPEPSKRLVDVLR-----VP-----HPDSQPDTHDHLQLLQSE-----RA 117
QY 264 GYSPPAKQLPGGRIHPSGYDND--RPNGLVRFCDV 297
Db 118 TLGPAVRPLPWQRV-----DRDVAPGTL-----CDV 143
RESULT 14
ELPG
pancreatic elastase (EC 3.4.21.36) I precursor - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 24-Apr-1984 #sequence_revision 30-Sep-1990 #text_change 09-Jul-2004

C;Accession: JS0013; A26777; A10061; A00959
R;Shirasu, Y.; Yoshida, H.; Miyakawa, T.; Matsuki, S.; Tanaka, J.I.; Ikenaga, H.
J. Biochem. 99, 1707-1712, 1986
A;Title: Isolation and expression in Escherichia coli of a cDNA clone encoding porcine p
A;Reference number: A92005; MUID:86304235; PMID:3528137
A;Accession: JS0013
A;Molecule type: mRNA
A;Residues: 1-266 <SHI>
A;Cross-references: UNIPROT:P00772; GB:X04036; GB:D00070; GB:N00070; NID:g1941; PIDN:CAA
R;Tani, T.; Kawashima, I.; Furukawa, H.; Ohmine, T.; Takiguchi, Y.
J. Biochem. 101, 591-599, 1987
A;Title: Characterization of a silent gene for human pancreatic elastase I: structure of
A;Reference number: A26777; MUID:87250343; PMID:3648024
A;Accession: A26777
A;Molecule type: mRNA
A;Residues: 1-125,'G',127-183,'L',185-266 <TAN>
A;Cross-references: GB:D00160; NID:g217683; PIDN:BA00118.1; PID:g217684
A;Note: the authors translated the codon GGG for residue 58 as Gln, GGC for residue 126 &
R;Shotton, D.M.; Hartley, B.S.
Biochem. J. 131, 643-675, 1973
A;Title: Evidence for the amino acid sequence of porcine pancreatic elastase.
A;Reference number: A90267; MUID:73229121; PMID:4578945
A;Accession: A10061
A;Molecule type: protein
A;Residues: 27-91,'N',93-203,'N',205-266 <SHO>
R;Shotton, D.M.; Hartley, B.S.
Nature 225, 811-816, 1970
A;Title: Three-dimensional structure of tosyl-elastase.
A;Reference number: A93160; MUID:70114044; PMID:5415110
A;Contents: annotation; X-ray crystallography, 3.5 angstroms; active site
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; pancreas; serine proteinase; zymogen
F;1-16/Domain: signal sequence #status predicted <SIG>
F;17-26/Domain: activation peptide #status predicted <APT>
F;27-266/Product: elastase I #status experimental <MAT>
F;27-259/Domain: trypsin homology <TRY>
F;56-72,153-220,184-200,210-240/Disulfide bonds: #status experimental
F;71,119,214/Active site: His, Asp, Ser #status experimental
Query Match 5.3%; Score 113; DB 1; Length 266;
Best Local Similarity 27.3%; Pred. No. 0.083;
Matches 36; Conservative 21; Mismatches 41; Indels 34; Gaps 6;
QY 147 NYPFSTSVKLSGTG-----CTGTLVAEKHVLTAACIHDGKTY--VKGPKLRVGFLLPKPK 198
Db 37 SMPQSISLQYRSGSSWAHTCGGTLIRQNNWMTAAACVDRELTFRVVVGHNH----- 88
QY 199 FKDGGGRANDTSAMPEQKQWIRVKTTHVPKGIKGNANDIGMDYDVALLEL-KKPKH 257
Db 89 -----NQNDGTE-----QYGVGVQIVVHPYW---NTDDVAAGYDIALRLAQSVTL 131
QY 258 RKFMKIGVSPPA 269
Db 132 NSYVLGVLPR 143
RESULT 15
ELRT1
pancreatic elastase (EC 3.4.21.36) I precursor - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 18-Aug-1982 #sequence_revision 18-Aug-1982 #text_change 09-Jul-2004
C;Accession: A00960; A20534
R;MacDonald, R.J.; Swift, G.H.; Quinto, C.; Swain, W.; Pictet, R.L.; Nikovits, W.; Rutter
Biochemistry 21, 1453-1463, 1982
A;Title: Primary structure of two distinct rat pancreatic preproelastases determined by
A;Reference number: A00960; MUID:82182967; PMID:6918221
A;Accession: A00960
A;Molecule type: mRNA
A;Residues: 1-266 <MAC>
A;Cross-references: UNIPROT:P00773; GB:V01234; NID:G56088; PIDN:CAA24544.1; PID:G56089
R;Largman, C.
Biochemistry 22, 3763-3770, 1983
A;Title: Isolation and characterization of rat pancreatic elastase.

Search completed: July 1, 2005, 21:09:19
Job time : 21.826 secs

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: July 1, 2005, 20:53:37 ; Search time 92.7095 Seconds
(without alignments)
2165.204 Million cell updates/sec

Title: US-09-658-677-15

Perfect score: 2131

Sequence: 1 MAGIPGLLLPFLLCVAGQ.....IKGNYLDCREGDVRPPGSN 392

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1612378 seqs, 512079187 residues

Total number of hits satisfying chosen parameters: 1612378

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Uniprot 03.*

1: uniprot_sprot.*

2: uniprot_trembl.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	2080	97.6	383	1 PS23 HUMAN	O95084 homo sapien
2	1903	89.3	383	2 O6AY61	O6AY61 rattus norv
3	1888.5	88.6	382	1 PS23 MOUSE	O96X66 mus muscucu
4	1882.5	88.3	382	2 O8BZS4	O8BZS4 mus muscucu
5	1042	48.9	413	2 O9BQF6	O9BQF6 homo sapien
6	1041	48.9	413	2 O8N320	O8N320 homo sapien
7	1016.5	47.7	409	2 O8COF9	O8COF9 mus muscucu
8	1014.5	47.6	418	2 O6GML6	O6GML6 brachydanio
9	1011.5	47.5	409	2 O8C0L5	O8C0L5 mus muscucu
10	1005.5	47.2	409	2 O8C0D6	O8C0D6 mus muscucu
11	194	9.1	316	1 GSEP BACLI	P80057 bacillus li
12	194	9.1	316	2 O65NR6	O65NR6 bacillus li
13	188	8.8	450	2 O71VE5	O71VE5 listeria mo
14	178.5	8.4	384	2 O98G17	O98G17 rhizobium l
15	158	7.4	271	2 O987W6	O987W6 rhizobium l
16	142.5	6.7	358	2 O931E7	O931E7 staphylococ
17	141.5	6.6	321	2 O7NGB4	O7NGB4 gloeobacter
18	138.5	6.5	323	2 O73D54	O73D54 bacillus ce
19	135.5	6.4	313	1 MPR BACSU	P39790 bacillus su
20	133.5	6.3	799	2 O6PF94	O6PF94 mus muscucu
21	133.5	6.3	811	1 TMS6 MOUSE	O811H5 bacillus ce
22	132.5	6.2	290	2 O81HL5	O81HL5 streptococ
23	131	6.1	218	2 O8DR41	O8DR41 streptococ
24	131	6.1	267	2 O6DGM4	O6DGM4 brachydanio
25	129.5	6.1	303	2 O9EXE9	O9EXE9 bacillus in
26	129	6.1	286	2 O6AZC0	O6AZC0 brachydanio
27	129	6.1	469	2 O9GMD9	O9GMD9 ornithorhyn
28	127.5	6.0	1322	2 O9NAT0	O9NAT0 anopheles g
29	126.5	5.9	678	2 O9JJ58	O9JJ58 rattus norv
30	124.5	5.8	1234	2 O7PIQ7	O7PIQ7 anopheles g
31	124.5	5.8	1322	2 O7Pnr7	O7Pnr7 anopheles g

32	124.5	5.8	1322	2	O9NJS5	O9NJS5 anopheles g
33	124	5.8	266	1	EL1 BOVIN	O28153 bos taurus
34	124	5.8	266	2	O46644	O46644 macaca fasc
35	123.5	5.8	339	2	O9QX91	O9QX91 rattus norv
36	123.5	5.8	366	2	O9QX85	O9QX85 rattus norv
37	123.5	5.8	541	2	O9QX90	O9QX90 rattus norv
38	123.5	5.8	623	2	O9QJ33	O9QJ33 rattus norv
39	123.5	5.8	643	2	O9QX84	O9QX84 rattus norv
40	122	5.7	259	2	O9XY61	O9XY61 ctencecephal
41	122	5.7	376	1	FA10_HOPST	P83370 hoptocephal
42	122	5.7	449	2	O9VDU8	O9VDU8 drosophila
43	121.5	5.7	490	1	FA10_RABIT	O19045 oryctolagus
44	121	5.7	745	2	O9QWF9	O9QWF9 triakis scy
45	121	5.7	1059	2	O7Z411	O7Z411 homo sapien

ALIGNMENTS

RESULT 1
PS23_HUMAN STANDARD; PRT; 383 AA.
AC O95084;
DT 28-FEB-2003 (Rel. 41, Created)
DT 28-FEB-2003 (Rel. 41, Last sequence update)
DT 25-OCT-2004 (Rel. 45, Last annotation update)
DE Serine protease 23 precursor (EC 3.4.21.-) (Putative secreted protein
DE ZSIG13) (UNQ270/PRO307).
GN Name=PRSS23; Synonyms=ZSIG13;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Euthera; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RC SEQUENCE FROM N.A.
RP TISSUE=Umbilical vein;
RA Li X., Tedder T.F.;
RT "A novel serine protease from human umbilical vein endothelial
RT cells.";
RL Submitted (JUL-1997) to the EMBL/GenBank/DBJ databases.
RN [2]
RC SEQUENCE FROM N.A.
RP Sheppard P., Blumberg H., Jelinek L., Foster D., O'Hara P.;
RL Submitted (OCT-1999) to the EMBL/GenBank/DBJ databases.
RN [3]
RC SEQUENCE FROM N.A.
RP TISSUE=Uterus;
RX MEDLINE=21154917; PubMed=11230166; DOI=10.1101/gr.154701;
RA Wiemann S., Weil B., Wellenreuther R., Gassenhuber J., Glassl S.,
Ansoerge W., Boecker M., Bloeker H., Bauersachs S., Blum H.,
Lauber J., Dueterhoff A., Beyer A., Koehrer K., Strack N.,
Mewes H.-W., Ottenwaelder B., Obermaier B., Tampe J., Heubner D.,
Wambutt R., Korn B., Klein M., Poustka A.;
RT "Towards a catalog of human genes and proteins: sequencing and
RT analysis of 500 novel complete protein coding human cDNAs.";
RL Genome Res. 11:422-435(2001).
RN [4]
RC SEQUENCE FROM N.A.
RP MEDLINE=22887296; PubMed=12975309; DOI=10.1101/gr.1293003;
RA Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,
Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,
Eaton D., Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S.,
Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,
Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,
Seshagiri S., Simmons L., Smith J., Singh J., Stinson J., Vagts A.,
Vandlen R., Watanabe C., Wieand D., Woods K., Xie M.-H., Yansura D.,
Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,
Godowski P., Gray A.;
RT "The secreted protein discovery initiative (SPDI), a large-scale
RT effort to identify novel human secreted and transmembrane proteins: a
RT bioinformatics assessment.";
RL Genome Res. 13:2265-2270(2003).
RN [5]

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RP SEQUENCE FROM N.A.
RC TISSUE=Cervix;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullaly S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettman M., Madan A., Young A.C., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Butterfield Y.S., Schein J.E., Jones S.J.M., Marra M.A.,
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.,
RT "Generation and initial analysis of more than 15,000 full-length human
and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
CC -!- SUBCELLULAR LOCATION: Secreted (Potential).
CC -!- SIMILARITY: Belongs to the peptidase S1 family.
-----
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CC between the Swiss Institute of Bioinformatics and the EMBL outstation -
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DR EMBL; AF015287; AD01553.1; -
DR EMBL; AF193611; AAF07186.1; -
DR EMBL; ALI36914; CAB66848.1; -
DR EMBL; AY359033; AAQ89392.1; -
DR EMBL; BC001278; AAH01278.1; -
DR MEROPS; S01.309; -
DR H-InvDB; HIX0010006; -
DR InterPro; IPR009003; Pept_Ser_Cys.
DR InterPro; IPR001254; Peptidase_S1.
DR InterPro; IPR001314; Peptidase_S1A.
DR Pfam; PF00089; Trypsin; 1.
DR PRINTS; PR00722; CHYMOTRYPSIN.
DR SMART; SM00020; Tryp_SPC; 1.
DR PROSITE; PS00240; TRYPSIN_DOM; FALSE_NEG.
DR PROSITE; PS00134; TRYPSIN_HIS; 1.
DR PROSITE; PS00135; TRYPSIN_SER; FALSE_NEG.
DR KMW Hydrolyase; Serine protease; Signal.
FT SIGNAL 1 23 Potential.
FT CHAIN 24 383 Serine protease 23.
FT ACT_SITE 175 175 Charge relay system (By similarity).
FT ACT_SITE 240 240 Charge relay system (By similarity).
FT ACT_SITE 316 316 Charge relay system (By similarity).
FT DISULFID 160 176 By similarity.
FT CARBOHYD 93 93 N-linked (GlcNAc...) (Potential).
FT CARBOHYD 207 207 N-linked (GlcNAc...) (Potential).
SQ SEQUENCE 383 AA; 43001 MW; 46E86C11ABFD558F CRC64;

Query Match 97.6%; Score 2080; DB 1; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.3e-167;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLFLFLPCLLCVAGQVSPYAPWPTWPAFLPVVLPOSTUNLAKPFGAEAKLE 60
DB 1 MAGIPGLFLFLPCLLCVAGQVSPYAPWPTWPAFLPVVLPOSTUNLAKPFGAEAKLE 60
QY 61 VSSSCGPGQCHKPTLYEAKQYLSYETLYANGSRTEQVGYIYLLSSGDAQHRDSCS 120
DB 61 VSSSCGPGQCHKPTLYEAKQYLSYETLYANGSRTEQVGYIYLLSSGDAQHRDSCS 120
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QY 121 SKSRRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGCTGTLVAEKHVLTAACHIHG 180
DB 121 SKSRRKQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGCTGTLVAEKHVLTAACHIHG 180
QY 181 KTYVKGTKLRVGFELKPKFKDGGRCANDSTSAPEQMKFQWIRVXKTHVPGKWIKNAND 240
DB 181 KTYVKGTKLRVGFELKPKFKDGGRCANDSTSAPEQMKFQWIRVXKTHVPGKWIKNAND 240
QY 241 IGWDYDYALLELKKPKKFKMIGVSPAKQIUPGGRHIFSGYDNDPRPGLNLYVRFCDVKDE 300
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QY 301 TYDLYQQCDAQPGAGSGGVYVMWKRQOQKWERKIIGIFSGHQWVMNGSPQDFNVAVR 360
DB 301 TYDLYQQCDAQPGAGSGGVYVMWKRQOQKWERKIIGIFSGHQWVMNGSPQDFNVAVR 360
QY 361 ITPLKYAQICYWIKGNLYDCREG 383
DB 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 2
Q6AY61 PRELIMINARY; PRT; 383 AA.
ID Q6AY61
AC Q6AY61;
DT 25-OCT-2004 (TReMBLrel. 28, Created)
DT 25-OCT-2004 (TReMBLrel. 28, Last sequence update)
DT 25-OCT-2004 (TReMBLrel. 28, Last annotation update)
DE Hypothetical protein.
OS Rattus norvegicus (Rat).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
OX NCBI_TaxID=10116;
RN [1]
RP SEQUENCE FROM N.A.
RX TISSUE=Kidney;
RC PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullaly S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettman M., Madan A., Young A.C., Rodrigues S., Sanchez A.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smallus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]
RP SEQUENCE FROM N.A.
RC TISSUE=Kidney;
RA Director MGC Project;
RL Submitted (AUG-2004) to the EMBL/GenBank/DBJ databases.
CC -!- SIMILARITY: Belongs to peptidase family S1.
DR EMBL; BC079179; AAH79179.1; -
DR GO; GO:0000786; C:nucleosome; IEA.
DR GO; GO:0005634; C:nucleus; IEA.
DR GO; GO:0004263; F:chymotrypsin activity; IEA.
DR GO; GO:0003677; F:DNA binding; IEA.
DR GO; GO:0007001; P:chromosome organization and biogenesis (sen. .; IEA.
DR GO; GO:0006334; P:chromosome assembly; IEA.
DR GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR InterPro; IPR001951; Histone_H4.
DR InterPro; IPR001254; Peptidase_S1.
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DR InterPro; IPR001314; Peptidase_S1A.
DR InterPro; IPR009003; Pept_Ser_Cys.
DR Pfam; PF00089; Trypsin; 1.
DR PRINTS; PR00722; CHYMOTRYPSIN.
DR PRODOM; PD01827; Histone H4; 1.
DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN 1.
KW Hydrolase; Hypothetical protein; Protease; Serine protease.
SQ SEQUENCE 383 AA; 41159 MW; 4AB12CC7B66CDFC8 CRC64;

Query Match 89.3%; Score 1903; DB 2; Length 383;
Best Local Similarity 91.4%; Pred. No. 1.3e-152;
Matches 350; Conservative 11; Mismatches 22; Indels 0; Gaps 0;

QY 1 MAGIPGLIFLLFELLCVAVGVSYSPKPKQTPAYRLPVVLPOSTINLAKPDGAEAKLE 60
DB 1 MAGIPGLIILLVLLCVFMQVSPVNPWKTPAYRLPIVLPOSTLKLAKPDPFADAKLE 60

QY 61 VSSSCGPOCHKGTPLPTYEAKQVLSYETLYANGSRRTQVGIYILSSSGDGAQHRS DGS 120
DB 61 VSSSCGPOCHKGTPLPTYEAKQVLSYETLYANGSRRTQVGIYILSSSGDGAQHRS DSEA 120

QY 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLTGCTGLVAEKHVLTAACHIDG 180
DB 121 AGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLTGCTGLVAEKHVLTAACHIDG 180

QY 181 KTVYVGTQKLVGFLPKPKFGKGRGANDSTAMPQKFWIRVKTTHVPKGIKGNAND 240
DB 181 KTVYVGTQKLVGFLPKPKFGKGRGANDSTAMPQKFWIRVKTTHVPKGIKGNAND 240

QY 241 IGMDDYDVALLEKPKPKFKMKGVSPPAKLPGRITHFSGYNDPGRNIVYRFDVKDE 300
DB 241 IGMDDYDVALLEKPKPKFKMKGVSPPAKLPGRITHFSGYNDPGRNIVYRFDVKDE 300

QY 301 TYDLLYQCCDAQPCAGSGGVYVWRMKRQQQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
DB 301 TYDLLYQCCDAQPCAGSGGVYVWRMKRQQQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360

QY 361 ITPKVAQICWIKGNVLDREG 383
DB 361 ITPKVAQICWIKGNVLDREG 383

RESULT 3

PS23_MOUSE STANDARD; PRT; 382 AA.
AC Q9D6X6; Q8VEG1;
DT 28-FEB-2003 (Rel. 41, Created)
DT 28-FEB-2003 (Rel. 41, Last sequence update)
DT 25-OCT-2004 (Rel. 45, Last annotation update)
DE Serine protease 23 precursor (EC 3.4.21.-).
GN Names=Prs23;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RS SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Tongue;
RX MEDLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;
RA Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,
RA Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,
RA Yagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori T.,
RA Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
RA Schriml L.M., Kanapin A., Matsuda H., Batalov S., Beisel K.W.,
RA Blake J.A., Bradt D., Brucci V., Chothia C., Corbani L.E., Cousins S.,
RA Dalla E., Dragani T.A., Fletcher C.F., Forrest A., Frazer K.S.,
RA Gaasterland T., Gariboldi M., Glessi C., Godzik A., Gough J.,
RA Grimmond S., Guetincich S., Hirokawa N., Jackson I.J., Jarvis E.D.,
RA Kanai A., Kawai H., Kawasawa Y., Kedzierzki R.M., King B.L.,
RA Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
RA Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,
RA Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,
RA Petrovskaya N., Pillai R., Pontius J.U., Qi D., Ramachandran S.,
RA Ravasi T., Reed J.C., Reed D.J., Reid J., Ring B.Z., Ringwald M.,
RA Sandelin A., Schneider C., Semple C.A., Setou M., Shimada K.,
RA Sultana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M.,
RA Verardo R., Wagner L., Wahlestedt C., Wang Y., Watanabe Y., Wells C.,
RA Wilming L.G., Wyndham-Boris A., Yanagisawa M., Yang I., Yang L.,
RA Yuan Z., Zavolan M., Zhu Y., Zimmer A., Carninci P., Hayatsu N.,
RA Hirozane-Klehtkawa T., Konno H., Nakamura M., Sakazume N., Sato K.,
RA Shiraki T., Waki K., Kawai J., Aizawa K., Arakawa T., Fukuda S.,
RA Hara A., Hasehizume W., Inotani K., Ishii Y., Itoh M., Kagawa I.,
RA Miyazaki A., Sakai K., Sasaki D., Shibata K., Shinagawa A.,
RA Yasunishi A., Yoshino M., Waterston R., Lander E.S., Rogers J.,
RA Birney E., Hayashizaki Y.;
RT "Analysis of the mouse transcriptome based on functional annotation of
RT 60,770 full-length cDNAs.";
RL Nature 420:563-573 (2002).
RN [2]
RP SEQUENCE FROM N.A.
RX TISSUE=Breast tumor;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.P., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ustin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S., Loquellano N.A., Peters G.J., Abramson R.D., Mullaly S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettman M., Madan A., Rodriguez S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalek U., Smalish D.E.,
RA Scherch A., Schein J.E., Jones S.J.M., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
CC -!- SUBCELLULAR LOCATION: Secreted (Potential).
CC -!- SIMILARITY: Belongs to the peptidase S1 family.
CC -----
CC This SWISS-PROT entry is copyright. It is produced through a collaboration
CC between the Swiss Institute of Bioinformatics and the EMBL outstation -
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CC or send an email to license@isb-sib.ch).
CC -----
CC EMBL; AK009847; BAB26541.1; -;
DR EMBL; AK078518; BAC37319.1; -;
DR EMBL; BC018517; AAH18517.1; -;
DR HSP; P00746; LDSU.
DR MEROPS; S01.309; -;
DR MGI; 1923703; 2310046G15Rik.
DR InterPro; IPR009003; Pept_Ser_Cys.
DR InterPro; IPR001254; Peptidase_S1.
DR InterPro; IPR001314; Peptidase_S1A.
DR Pfam; PF00089; Trypsin; 1.
DR PRINTS; PR00722; CHYMOTRYPSIN.
DR SMART; SM00020; TRYP_SPC; 1.
DR PROSITE; PS00240; TRYPSIN_DOM; FALSE_NEG.
DR PROSITE; PS00134; TRYPSIN_HIS; 1.
DR PROSITE; PS00135; TRYPSIN_SER; FALSE_NEG.
KW Hydrolase; Serine protease; Signal.
FT SIGNAL 1 22 Potential.
FT CHAIN 23 382 Serine protease 23.
FT ACT_SITE 174 174 Charge relay system (By similarity).
FT ACT_SITE 239 239 Charge relay system (By similarity).
FT ACT_SITE 315 315 Charge relay system (By similarity).
FT DISULFID 159 175 By similarity.
FT CARBOHYD 92 N-linked (GlcNAc...) (Potential).

FT	CARBOHYD	206	206	N-linked (GlcNAc...)	(Potential).
FT	CONFLICT	260	260	M -> I (in Ref. 1; BAB26541).	
SQ	SEQUENCE	382 AA;	43071 MW;	6F09A5C80A5B2306 CRC64;	
Query Match					
Best Local Similarity 88.6%; Score 1888.5; DB 1; Length 382;					
Matches 348; Conservative 14; Mismatches 20; Indels 1; Gaps 1;					
QY	1	MAGIPGLLFLFLLCVAGQVSPYAPWKPTWPAYRLPVVLPQSTLNLAQDPFGABAKLE	60		
DB	1	MAGIPG-LFILLVLLCVFMQVSPYTPWKPTWPAYRLPVVLPQSTLNLAQDPFGABAKLE	59		
QY	61	VSSCGPQCHKGTPLTPTVEEAKQYLSYETLYANGSRTETQVGIYILSSGCGAQRDSGS	120		
DB	60	VSSCGPQCHKGTPLTPTVEEAKQYLSYETLYANGSRTETQVGIYILSSGCGAQRDSGA	119		
QY	121	SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG	180		
DB	120	TGSRKRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG	179		
QY	181	KTYVKGTPQKLRVGLFKPKFQKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND	240		
DB	180	KTYVKGTPQKLRVGLFKPKFQKDGGRGANDSTSSAMPDKMKFQWIRVKRTHVPKGIKNAND	239		
QY	241	IGMDYDYLLELKKPHKRFPMKIGVSPPAKQLPGGRIFHSFGYNDNDRPGLVYRFDVKDE	300		
DB	240	IGMDYDYLLELKKPHKRFPMKIGVSPPAKQLPGGRIFHSFGYNDNDRPGLVYRFDVKDE	299		
QY	301	TYDLLYQCCDAQPGASGSGVYVWMKRCQKQKWKIIGIFPSGHWDMNGSPQDENVAVR	360		
DB	300	TYDLLYQCCDAQPGASGSGVYVWMKRCQKQKWKIIGIFPSGHWDMNGSPQDENVAVR	359		
QY	361	ITPLKYAQICYWIKGNLYDCREG 383			
DB	360	ITPLKYAQICYWIKGNLYDCREG 382			
RESULT 4					
Q8BZS4	PRELIMINARY; PRT; 382 AA.				
AC	Q8BZS4				
DT	01-MAR-2003 (TEMBLrel. 23, Created)				
DT	01-MAR-2003 (TEMBLrel. 23, Last sequence update)				
DE	Mus musculus adult male cecum cDNA, RIKEN full-length enriched				
DE	library, clone:9130215B18 product:SERINE PROTEASE (HYPOTHETICAL 43.0				
DE	kDa PROTEIN) (PROTEASE, SERINE, 23) homolog.				
OC	Mus musculus (Mouse)				
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;				
OC	Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.				
OX	NCBI_TaxID=10090;				
RN	[1]				
RP	SEQUENCE FROM N.A.				
RC	STRAIN=C57BL/6J; TISSUE=Cecum;				
RX	MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;				
RA	Carninci P., Hayashizaki Y.;				
RT	"High-efficiency full-length cDNA cloning.;"				
RL	Meth. Enzymol. 303:19-44(1999).				
RN	[2]				
RP	SEQUENCE FROM N.A.				
RC	STRAIN=C57BL/6J; TISSUE=Cecum;				
RX	MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;				
RA	RIKEN FANTOM Consortium;				
RT	"Functional annotation of a full-length mouse cDNA collection.;"				
RL	Nature 409:685-690(2001).				
RN	[3]				
RP	SEQUENCE FROM N.A.				
RC	STRAIN=C57BL/6J; TISSUE=Cecum;				
RA	The FANTOM Consortium;				
RT	"Analysis of the mouse transcriptome based on functional annotation of				
RT	60,770 full-length cDNAs.;"				
RL	Nature 420:563-573(2002).				

RN	[4]	SEQUENCE FROM N.A.			
RC	STRAIN=C57BL/6J; TISSUE=Cecum;				
RX	MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;				
RA	Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K., Itoh M.,				
RA	Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;				
RT	"Normalization and subtraction of cap-trapper-selected cDNAs to				
RL	prepare full-length cDNA libraries for rapid discovery of new genes.;"				
RL	Genome Res. 10:1617-1630(2000).				
RN	[5]	SEQUENCE FROM N.A.			
RC	STRAIN=C57BL/6J; TISSUE=Cecum;				
RX	MEDLINE=20530913; PubMed=11076861; DOI=10.1101/gr.152600;				
RA	Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P.,				
RA	Konno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itoh M.,				
RA	Sumi N., Ishii Y., Nakamura S., Hazama M., Nishine T., Harada A.,				
RA	Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,				
RA	Fujisawa S., Inoue K., Togawa Y., Izawa M., Ohara E., Watahiki M.,				
RA	Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsura S., Kawai J.,				
RA	Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.;				
RT	"RIKEN integrated sequence analysis (RISA) system-384-format				
RL	sequencing pipeline with 384 multicapillary sequencer.;"				
RL	Genome Res. 10:1757-1771(2000).				
RN	[6]	SEQUENCE FROM N.A.			
RC	STRAIN=C57BL/6J; TISSUE=Cecum;				
RA	Adachi J., Aizawa K., Akimura T., Arakawa T., Bono H., Carninci P.,				
RA	Fukuda S., Furuno M., Hasegaki T., Hara A., Hashizume W.,				
RA	Hayashida K., Hayatsu N., Hiramoto K., Hiraoka T., Hirozane T.,				
RA	Hori F., Imotani K., Ishii Y., Itoh M., Kagawa I., Kasukawa T.,				
RA	Kurihara C., Matsuyama T., Miyazaki A., Murata M., Nakamura M.,				
RA	Nishi K., Nomura K., Numazaki R., Ohno M., Ohsato N., Okazaki Y.,				
RA	Saito R., Saitoh H., Sakai C., Sakai K., Sakazume N., Sano H.,				
RA	Sasaki D., Shibata K., Shinagawa A., Shiraki T., Sogabe Y., Tagami M.,				
RA	Tagawa A., Takahashi F., Takaku-Akai H. S., Takeda Y., Tanaka T.,				
RA	Tomaru A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y.;				
RL	Submitted (JUL-2001) to the EMBL/GenBank/DBJ databases.				
CC	-!- SIMILARITY: Belongs to peptidase family S1.				
DR	EMBL; AK033671; BAC28420.1; -				
DR	GO; GO:0004263; F:chymotrypsin activity; IEA.				
DR	GO; GO:0008233; F:peptidase activity; IEA.				
DR	GO; GO:0004295; F:trypsin activity; IEA.				
DR	GO; GO:0006508; P:proteolysis and peptidolysis; IEA.				
DR	InterPro; IPR001254; Peptidase S1.				
DR	InterPro; IPR001314; Peptidase S1A.				
DR	InterPro; IPR009003; Pept Ser_Cys.				
DR	PRINTS; PR00722; CHYMOTRYPSIN.				
DR	SMART; SM00020; Tryp_SPC; 1.				
DR	PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.				
KW	Hydrolase; Hypothetical protein; Protease; Serine protease.				
SQ	SEQUENCE 382 AA; 43147 MW; 556789818E12A081 CRC64;				
Query Match 88.3%; Score 1882.5; DB 2; Length 382;					
Best Local Similarity 90.6%; Pred. No. 6.8e-151;					
Matches 347; Conservative 14; Mismatches 21; Indels 1; Gaps 1;					
QY	1	MAGIPGLLFLFLLCVAGQVSPYAPWKPTWPAYRLPVVLPQSTLNLAQDPFGABAKLE	60		
DB	1	MAGIPG-LFILLVLLCVFMQVSPYTPWKPTWPAYRLPVVLPQSTLNLAQDPFGABAKLE	59		
QY	61	VSSCGPQCHKGTPLTPTVEEAKQYLSYETLYANGSRTETQVGIYILSSGCGAQRDSGS	120		
DB	60	VSSCGPQCHKGTPLTPTVEEAKQYLSYETLYANGSRTETQVGIYILSSGCGAQRDSGA	119		
QY	121	SGKSRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG	180		
DB	120	TGSRKRKRQIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG	179		
QY	181	KTYVKGTPQKLRVGLFKPKFQKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKNAND	240		
DB	180	KTYVKGTPQKLRVGLFKPKFQKDGGRGANDSSAMPDKMKFQWIRVKRTHVPKGIKNAND	239		

QY 241 IGMDDYDYLLEKPKHKKFPMKIGVSPAPKQLPGRHIFSGYDNDPGLNLYRFCVCKDE 300
 DB 240 IGMDDYDYLLEKPKHKKFPMKIGVSPAPKQLPGRHIFSGYDNDPGLNLYRFCVCKDE 299
 QY 301 TYDLYLQCCDAQPGASGSGVYVWRKQKQKWKIIGIPSGHQMNGSPQDFNVAVR 360
 DB 300 TYDLYLQCCDAQPGASGSGVYVWRKQKQKWKIIGIPSGHQMNGSPQDFNVAVR 359
 QY 361 ITPKYAQICWIKGNLYDCREG 383
 DB 360 ITPKYAQICWIKGNLYDCREG 382

RESULT 5
 Q9BQ6 PRELIMINARY; PRT; 413 AA.
 ID Q9BQ6
 AC Q9BQ6; 17, Created
 DT 01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
 DT 01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
 DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
 DE D1223E3.1 (Putative secreted protein ZS1G13).
 GN Names=dJ223E3.1;
 OS Homo sapiens (Human).
 OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
 OC Mammalia; Euthera; Primates; Catarrhini; Hominidae; Homo.
 OX NCBI_TaxID=9606;
 RN [1]
 RP SEQUENCE FROM N.A.
 RA Dunn M.;
 RL Submitted (MAR-2001) to the EMBL/GenBank/DBJ databases.
 RC -1- SIMILARITY: Belongs to peptidase family S1.
 DR EMBL; AL121939; CAC35071.1; -;
 DR MEROPS; S01.994; -; PRSS35.
 DR Genew; HGNC:21387; PRSS35.
 DR GO; GO:0004263; F:chymotrypsin activity; IEA.
 DR GO; GO:0008233; F:peptidase activity; IEA.
 DR GO; GO:0004295; F:trypsin activity; IEA.
 DR GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
 DR InterPro; IPR01254; Peptidase S1.
 DR InterPro; IPR01314; Peptidase S1A.
 DR InterPro; IPR009003; Pept Ser_Cys.
 DR Pfam; PF00089; Trypsin; 1.
 DR PRINTS; PR00722; CHYMOTRYPSIN.
 DR SMART; SM00020; Tryp_SPC; 1.
 DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
 DR Hydrolase; Protease; Serine protease.
 KW SEQUENCE 413 AA; 47098 MW; 818D9C951BD2D6C1 CRC64;

Query Match 48.9%; Score 1042; DB 2; Length 413;
 Best Local Similarity 47.8%; Pred. No. 1.1e-79;
 Matches 200; Conservative 68; Mismatches 100; Indels 50; Gaps 7;

QY 7 LFLFLFLLCAVG--QVSPYAPKPTWPAAYLPVLPVLPSTLNLAKPFGAEAKLEVSSS 64
 DB 5 LLWLIF--TPGWTLDGSEMDPFWHLKVPYPRIVSERTFHLTSPAEADAKMNV 61
 QY 65 CGPQCHGTPLPYERAKQVLSYETLYANGSRFTQVGI--YLLSSGCGAHRDSSGSG 122
 DB 62 CGTECQKELPTSLSELDYLYETVFENGTRTLTKVQDLVLEPT-----QNITTKG 115
 QY 123 KS--RRKQYGVDSRESIFGKDLLNYPSTSVKLSGTCTGLVAEKHVLTAHCIDCK 181
 DB 116 VSVRRKQVGTDSRESILDKRLTFNPFSTAVKLSGTGGLISQHVLTAAHCVDHCK 175
 QY 182 TVYKGTQKLVGPKFKDQGR-----GANDSTSAHPQMK----- 218
 DB 176 DYVKGSKLVGLLKNKSGGKRGSKRSREASGGQRETRHLERAKGRRRK 235
 QY 219 -----FQIRVKTTHVPKGIWTKGNANDIGMDYDYLLEKPKHKKFPMKIGV 265
 DB 236 SGRGQRIAEGRPSFQWTRVKNTHPIKGWARGMGDGLDYDYLLEKPKHKKYKWLGI 295
 QY 266 SPPAKQLPGRHIFSGYDNDPGLNLYRFCVCKDEYDLYLQCCDAQPGASGSGVYVWRM 325

DB 296 SPTIKKMPGMIHFSGFNDRADQLVYRFSVSDSNLLQYCDAESGSGTVYLRK 355
 QY 326 KQOQKWKIIGIPSGHQMNGSPQDFNVAVRITPLKYAQICWIKGNLYDCREG 383
 DB 356 DPDKKWKIIGIPSGHQMNGSPQDFNVAVRITPLKYAQICWIKGNLYDCREG 413

RESULT 6
 Q9N320 PRELIMINARY; PRT; 413 AA.
 ID Q9N320
 AC Q9N320;
 DT 01-OCT-2002 (TrEMBLrel. 22, Created)
 DT 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
 DT 25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
 DE Hypothetical protein PRSS35 (ENML522).
 GN Name=PRSS35; ORFNames=UNQ522;
 OS Homo sapiens (Human).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Mammalia; Euthera; Primates; Catarrhini; Hominidae; Homo.
 OX NCBI_TaxID=9606;
 RN [1]
 RP SEQUENCE FROM N.A.
 RA TISSUE=Brain;
 RC MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
 RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
 RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
 RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
 RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
 RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
 RA Brownstein M.J., Udwin T.B., Toshiyuki S., Carninci P., Prange C.,
 RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullany S.J.,
 RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
 RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
 RA Fahey J., Helton E., Kettman M., Madan A., Rodriguez S., Sanchez A.,
 RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
 RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
 RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
 RA Krzywinski M.I., Skalska U., Smallos D.E., Schnerch A., Schein J.E.,
 RA Jones S.J., Marra M.A.;
 RT "Generation and initial analysis of more than 15,000 full-length human
 and mouse cDNA sequences.";
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
 RN [2]
 RP SEQUENCE FROM N.A.
 RA TISSUE=Brain;
 RC Strausberg R.;
 RL Submitted (SEP-2002) to the EMBL/GenBank/DBJ databases.
 RN [3]
 RP SEQUENCE FROM N.A.
 RX MEDLINE=22987296; PubMed=12975309; DOI=10.1101/gr.1293003;
 RA Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,
 RA Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,
 RA Eaton D., Foster J., Grimaldi C., Gu Q., Hase P.E., Heldens S.,
 RA Huang A., Kim H.S., Klinkowski L., Jin Y., Johnson S., Lee J.,
 RA Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,
 RA Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,
 RA Vandlen R., Watanabe C., Wieand D., Woods K., Xie M.H., Yansura D.,
 RA Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,
 RA Godowski P.;
 RT "The secreted protein discovery initiative (SPDI), a large-scale
 effort to identify novel human secreted and transmembrane proteins: a
 bioinformatics assessment.";
 RL Genome Res. 13:2265-2270(2003).
 CC -1- SIMILARITY: Belongs to peptidase family S1.
 DR EMBL; BC037170; AAH37170.1; -;
 DR EMBL; AY358661; AAQ89024.1; -;
 DR GO; GO:0004263; F:chymotrypsin activity; IEA.
 DR GO; GO:0008233; F:peptidase activity; IEA.
 DR GO; GO:0004295; F:trypsin activity; IEA.
 DR GO; GO:0006508; P:proteolysis and peptidolysis; IEA.

RL Submitted (JUN-2004) to the EMBL/GenBank/DBJ databases.

CC -1- SIMILARITY: Belongs to peptidase family S1.
 DR EMBL; AK031411; BAC27392.1; -.
 DR EMBL; BC075675; AAH75675.1; -.
 DR MGD; MGI:2444800; Prs935.
 DR GO; GO:0005615; C:extracellular space; TAS.
 DR InterPro; IPR001254; Peptidase_S1.
 DR InterPro; IPR001314; Peptidase_S1A.
 DR InterPro; IPR009003; Pept_Ser_Cys.
 DR Pfam; PF00089; Trypsin; 1.
 DR PRINTS; PR00722; CHYMOTRYPSIN.
 DR SMART; SM00020; Tryp_SPC; 1.
 DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
 DR Hydrolase; Protease; Serine protease.
 SQ SEQUENCE 409 AA; 45787 MW; 5E22D4A908E7EPE5 CRC64;

Query Match 47.7%; Score 1016.5; DB 2; Length 409;
 Best Local Similarity 49.4%; Pred. No. 1.6e-77;
 Matches 193; Conservative 68; Mismatches 83; Indels 47; Gaps 5;
 QY 31 TWPAYRLPVVLPVLTNLAKEPFGAEAKLEVSSECGPCHGKGTPLTYEAKQYLSYETL 90
 DB 28 TWHLRIPOVQVSENTIHLASPTFOAGVVKATVCGIEQOEELPAPSLQLESLSYETI 87
 QY 91 YANGSRTEQVGYIL-----SSSGDGAQHRDSSGSKRRKQIYGYDSRFSIFGKDF 144
 DB 88 FENGTHLIRVQGLVLETRNSVKGAI-----PRRRQVGTDSRFSILDKRF 138
 QY 145 LLNYPFSTSVKLTGCTGTVAEKHVLTAACHDGTYYKGTQKLRVGLPKPKFDG-- 202
 DB 139 ATNFPNTAVKLTGCGTLVSPNHVLTAAHCVDHGDYVKGSKLVRVGLKWRNKGGRK 198
 QY 203 -GRGA-----NDSTSAMPEQM-----KFWIRVRKTHVPKG 232
 DB 199 KRGSKRSRRREASAGSOAHLRETTQREGKSRGRPRVTQGRPSQWTRVAKSTHPKG 258
 QY 233 WIKGNANDIGMDYDVALLELKKPKRKFMIKIGVSPPAKQLPGGRIHFSGYDNDPRGNLVY 292
 DB 259 WVRGEGGLALDYDVALLELKKRAHQHMLGVSPTITKLPGRRIHFSGDNDRDQLVY 318
 QY 293 RFCDVKDETVDLLYQQCDAQGASGVYVVRMKRQKQKWKRIIGFSGHGWDMNGSP 352
 DB 319 RFCSVSEESNDLLYQCYDAEAGSTGSGIYLRLEKPGQKWKRIVAVYSGHQWVDVHGQ 378
 QY 353 QDNVAVRITPLKYAQICVWIKGNYLDCREG 383
 DB 379 KDYNVAVRITPLKYAQICLWINGNAANCAYG 409

RESULT 8
 Q6GML6
 ID Q6GML6 PRELIMINARY; PRT; 418 AA.
 AC Q6GML6;
 DT 05-JUL-2004 (TEMBLrel. 27, Created)
 DT 05-JUL-2004 (TEMBLrel. 27, Last sequence update)
 DT 05-JUL-2004 (TEMBLrel. 27, Last annotation update)
 DE Zgc:91804.
 GN ORFNames=zgc:91804;
 OS Brachydanio rerio (Zebrafish) (Danio rerio).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
 OC Cyprinidae; Danio.
 OX NCBI_TaxID=7955;
 RN [1]
 RP SEQUENCE FROM N.A.
 RC TISSUE=Whole;
 RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
 RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
 RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
 RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
 RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
 RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
 RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullany S.J.,
 RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.O., Hulyk S.W.,
 RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
 RA Fahey J., Helton E., Kettman M., Madan A., Rodriguez S., Sanchez A.,
 RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
 RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
 RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
 RA Krzywinski M.I., Skalska U., Smallos D.E., Schnerch A., Schein J.E.,
 RA Jones S.J., Marra M.A.;
 RT "Generation and initial analysis of more than 15,000 full-length human
 and mouse cDNA sequences.";
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
 RN [2]
 RP SEQUENCE FROM N.A.
 RC TISSUE=Whole;
 RA Strausberg R.;
 RL Submitted (JUN-2004) to the EMBL/GenBank/DBJ databases.
 DR EMBL; BC074028; AAH74028.1; -.
 DR ZFIN; ZDB-GENE-040704-55; zgc:91804.
 DR GO; GO:0008233; F:peptidase activity; IEA.
 DR GO; GO:0004295; F:trypsin activity; IEA.
 DR GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
 DR InterPro; IPR001254; Peptidase_S1.
 DR InterPro; IPR009003; Pept_Ser_Cys.
 DR Pfam; PF00089; Trypsin; 1.
 DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
 DR Hydrolase; Protease; Serine protease.
 SQ SEQUENCE 418 AA; 47270 MW; AF53345CEB94F720 CRC64;
 Query Match 47.6%; Score 1014.5; DB 2; Length 418;
 Best Local Similarity 46.6%; Pred. No. 2.4e-77;
 Matches 196; Conservative 64; Mismatches 118; Indels 43; Gaps 2;
 QY 1 MAGIPCLLFLFLCAVGVSPYSAPMKPTPAVRLPVVLPVLTNLAKEPFGAKLE 60
 DB 1 MGPVPLTLLLSALAVLSTTVDPTDGTTPQKIPLVQEKQTVHLSSEFLAKPND 60
 QY 61 VSSSCGPQCHGTPLPTVEEAKQYLSYETLYANGSRTEQVGIYILSSGSGDAQHRDSSG 120
 DB 61 LHGICGIECQQLPEPSLDLLEQLLSYETMYDNGTRTLTTVTQDLNVSNDWT---GAS 116
 QY 121 SKSRKQRIYGYDSRFSIFGKDFLLNVFPSTSVKLTGCTGTVAEKHVLTAACHIDG 180
 DB 117 QLHTRHREVYGTDRFTITDKOYSMKYFPSTSVKISTGCGVLVSPKHLVTAACHIHNG 176
 QY 181 KTYVKGTKLRVGLKP-----KFKD 201
 DB 177 TTYLGVQLSVGLKRSRRKNGRKGKRGKQKHEEVDENGEIVEKQERKSKG 236
 QY 202 GGRGANDSTSAMPEQMKFQWIRVRKTHVPKGIKGNANDIGMDYDVALLELKKPKRKF 261
 DB 237 KGRNRSRSTDSQSPSRWTRVKQVPGKGFISENVLADYDVALLEKRAQTKFM 296
 QY 262 KIGVSPPAKQLPGGRIHFSGYDNDPRGNLVYPCDVKDETVDLLYQQCDAQGASGVY 321
 DB 297 DLGVIPSVKKLPAGRIFHFSGFDPRGNLVYRFSVSEESNDLLYQCYDAKPGSSGVY 356
 QY 322 VVMKRRQKQKWKRIIGFSGHGWDMNGSPDENVAVRITPLKYAQICVWIKGNVLDGR 381
 DB 357 IRLKPGKKKKWKRIIGVFSGHQWVDVNGQQQDYNVAVRITPLKYAQICRWVHGDSQCR 416
 QY 382 E 382
 DB 417 D 417
 RESULT 9
 Q8COL5
 ID Q8COL5 PRELIMINARY; PRT; 409 AA.
 AC Q8COL5;
 DT 01-MAR-2003 (TEMBLrel. 23, Created)

3] SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Testis;
RA The PANTOM Consortium;
RT "Analysis of the mouse transcriptome based on functional annotation of
RL 60,770 full-length cDNAs";
RN Nature 420:563-573(2002).
4] SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Testis;
RX MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;
RA Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K., Itoh M.,
RA Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;
RT "Normalization and subtraction of cap-trapper-selected cDNAs to
RL prepare full-length cDNA libraries for rapid discovery of new genes";
RN Genome Res. 10:1617-1630(2000).
5] SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Testis;
RX MEDLINE=20530913; PubMed=11076861; DOI=10.1101/gr.152600;
RA Shibata K., Itoh M., Aizawa K., Nagaoaka S., Sasaki N., Carninci P.,
RA Konno H., Akiyama J., Nishi K., Kitsuina T., Tashiro H., Itoh M.,
RA Sumi N., Ishii Y., Nakamura S., Hazama M., Nishine T., Harada A.,
RA Yamamoto R., Matsuoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,
RA Fujiwaka S., Inoue K., Togawa Y., Izawa M., Ohara E., Watabiki M.,
RA Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,
RA Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.;
RT "Riken integrated sequence analysis (RISA) system-384-format
RL sequencing pipeline with 384 multicapillary sequencer";
RN Genome Res. 10:1757-1771(2000).
6] SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Testis;
RA Adachi J., Aizawa K., Akimura T., Arakawa T., Bono H., Carninci P.,
RA Fukuda S., Furuno M., Hanganaki T., Hara A., Hashizume W.,
RA Hayashida K., Hayatsu N., Hiramoto K., Hiraoka T., Hirozane T.,
RA Hori F., Imotani K., Ishii Y., Itoh M., Kagawa I., Kasukawa T.,
RA Katoh H., Kawai J., Kojima Y., Kondo S., Konno H., Kouda M., Koya S.,
RA Kurihara C., Matsuyama T., Miyazaki A., Murata M., Nakamura M.,
RA Nishi K., Nomura K., Numazaki R., Ono M., Ohsato N., Okazaki Y.,
RA Saito R., Saitoh H., Sakai C., Sakai K., Sakazume N., Sano H.,
RA Saeki D., Shibata K., Shinagawa A., Shiraki T., Sogabe Y., Tagami M.,
RA Tagawa A., Takahashi F., Takaku-Akai H., Tanaka Y., Tanaka I.,
RA Tomaru A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y.;
RL Submitted (JUL-2001) to the EMBL/GenBank/DBJ databases.
DR EMBL; AK031644; BAC27491.1; -;
DR MGD; MGI:2444800; P8a35.
DR GO; GO:0005615; C:extracellular space; TAS.
DR InterPro; IPR001254; Peptidase_S1.
DR InterPro; IPR009003; Pept_Ser_Cys.
DR Pfam; PF00089; Trypsin; 1.
DR SMART; SM00020; Tryp_Spc; 1.
DR Hydrolase; Protease; Serine protease.
KW SEQUENCE 409 AA; 45736 MW; E9C9B91B6800719C CRC64;
SQ
Query Match 47.2%; Score 1005.5; DB 2; Length 409;
Best Local Similarity 48.8%; Pred. No. 1.4e-76;
Matches 191; Conservative 70; Mismatches 83; Indels 47; Gaps 5;
QY 31 TWPAYRLPVLPQSTLNLAKEPDEGAELKLVSSSCGQCHKGTPLPYEEAKQYLSVETL 90
DB 28 TWHLRSIPQVSVTHLHASPFTQADAGVVKATVCGIECQEEPLPAPLSQLESLSYETI 87
QY 91 YANGSRTEQTQVGIYL-----SSSGDGAQHRSKSSGSKRRKQIYGYDSRFSIFKQDF 144
DB 88 FENGTRTLTRVKQGLVLETRNSSVKGAH-----PRRRQVGTDSRFSILDKRF 138
QY 145 LLNYPFSTSVKLSGCTGTLVAEKHVLTAACHDGYKTKYVKGQKLRVGLKPKFKDG-- 202
DB 139 ATNFPFNTAVKLSGCTGLVSPNHLTAANCVDHGDYKVGSKLAVGLKVRNGGRK 198
QY 203 -GRGA-----NDSTSAMPEQM-----KQWIRVVRTHVPKG 232

DB 199 KRGSKSRREASAGOSQAHLESTTORPGKSRRCPRVYTOGRPPSFQWTRVSKTHIPKG 258
QY 233 WIKGNANDIGMDYDVALLELKKPKRPMKIGVSPPAKQIPGGRHIFSGYDNDRPGNLVY 292
DB 259 WVRGEGGLADYDVALLELKKRAHQHQMELGVSPTITKLPGGQIHFSGFNDRDEQLVY 318
QY 293 RFGDVKDEYDLYOCCDAPGASGGVYVMKQOQKWKRTIIGFSGHGVMDNGSP 352
DB 319 RFGSVSEESNDLLYQYCDAAEGSTGSGIYLRLEKPGQKNWKRKIVAVYSGHGVMDVHVGQ 378
QY 353 QDFNVAVRITPLKYAQICVYKNGNYLDCREG 383
DB 379 KDYNVAVRITPLKYAQICLWIHGNAANCAGV 409
RESULT 11
GSEP_BACLI STANDARD; PRT; 316 AA.
AC P80057;
DT 01-MAY-1992 (Rel. 22, Created)
DT 01-OCT-1993 (Rel. 27, Last sequence update)
DT 25-OCT-2004 (Rel. 45, Last annotation update)
DE Glutamyl endopeptidase precursor (SC 3.4.21.19) (Glutamate specific
DE endopeptidase) (GSE).
GN Name:blase;
OS Bacillus licheniformis.
OC Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus.
OX NCBI_TaxID=1402;
RN [1]
RP SEQUENCE FROM N.A., AND PARTIAL SEQUENCE.
RC STRAIN=ATCC 14580;
RX MEDLINE=93054737; PubMed=1429718;
RA Kakudo S., Kikuchi N., Kitadokoro K., Fujiwara T., Nakamura E.,
RA Okamoto H., Shin M., Tamaki M., Teraoka H., Tezuka H., Yoshida N.;
RT "Purification, characterization, cloning, and expression of a glutamic
RT acid-specific protease from Bacillus licheniformis ATCC 14580";
RL J. Biol. Chem. 267:23782-23788(1992).
RN [2]
RP SEQUENCE OF 95-316.
RX MEDLINE=92155199; PubMed=1346764;
RA Svendsen I., Bredam K.;
RT "Isolation and amino acid sequence of a glutamic acid specific
RT endopeptidase from Bacillus licheniformis";
RL Eur. J. Biochem. 204:165-171(1992).
CC -1- FUNCTION: Specific for hydrolysis of peptide bonds on the carboxyl
CC side of acidic amino acid residues, with a strong preference for
CC Glu.
CC -1- CATALYTIC ACTIVITY: Preferential cleavage: Asp-|-Xaa, Glu-|-Xaa.
CC -1- SUBCELLULAR LOCATION: Secreted.
CC -1- SIMILARITY: Belongs to the peptidase S1B family.
CC
CC This SWISS-PROT entry is copyright. It is produced through a collaboration
CC between the Swiss Institute of Bioinformatics and the EMBL Outstation -
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CC entities requires a license agreement (See <http://www.isb-sib.ch/announce/>
CC or send an email to license@isb-sib.ch).
CC
CC EMBL; D10060; BAA00949.1; -;
DR PIR; A45134; A45134.
DR MEROPS; S01.271; -;
DR InterPro; IPR000126; Pept_S1B_AS.
DR InterPro; IPR009003; Pept_Ser_Cys.
DR InterPro; IPR001254; Peptidase_S1.
DR InterPro; IPR008256; Peptidase_S1B.
DR Pfam; PF00089; Trypsin; 1.
DR PRINTS; PR00839; V8PROTEASE.
DR SMART; SM00020; Tryp_Spc; 1.
DR PROSITE; PS00672; V8_HIS; 1.
DR PROSITE; PS00673; V8_SER; 1.
KW Direct protein sequencing; Hydrolase; Serine protease; Signal.


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Db      169  YGSCSNTLIGPRTVLTAHCLYSHEDKDWL---SEYLFV-----PGLNGSTA--- 212
Qy      215  EQMKFQIWRVAKRTHVPKGIWKNANDIG--MDYDYALLEKKPKHRKFMKIGVSPPAKOL 272
Db      213  DDAPFGAFTYESAYVLQGFIDNYQGYGVSIPWDLGIITLKQDVCTNLGWLGYA-NYDDL 271
Qy      273  PGGRIHFSGYNDNRP--GNLVYRFCDKDETVDLLYQQ--CDAQPGASGSGVYVRMWKROQ 329
Db      272  GDFETANLVGPGDKPMGTMKWCASEVHAENIAPYFYDCTDTPFGSSGSSVAYDTKSKQ 331
Qy      330  QKWERKIIFSGHQWDMNGSPQDFNVAVRITPLKYAQICVWIKGNY 377
Db      332  R-----IITG---VNVASEP-DANTAVRLN-----AANQVWINSLY 363

RESULT 15
Q987W6
ID Q987W6 PRELIMINARY; PRT; 271 AA.
AC Q987W6;
DT 01-OCT-2001 (TRENBLrel. 18, Created)
DT 01-OCT-2001 (TRENBLrel. 18, Last sequence update)
DT 01-MAR-2004 (TRENBLrel. 26, Last annotation update)
DE Glutamic acid specific endopeptidase.

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[illegible]

Qy 324 MWKQOQKWKIIGI-----FSGHOWDMNGSPD-----FNVAVRITP 363
Db 210 -WIHQAGPPVVIAGPRPHSGGAWGCRPGVPLAPAGLEFNRGVRLTP 259

Search completed: July 1, 2005, 21:08:12
Job time : 94.7095 secs

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: July 1, 2005, 20:52:53 ; Search time 98.1294 Seconds
(without alignments)
1509.530 Million cell updates/sec

Title: US-09-658-677-18

Perfect score: 2080

Sequence: 1 MAGIPGLFLFLFLCAVGQ.....LKVAICYWIKGNVLDREG 383

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 2105692 seqs, 386760381 residues

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : A Geneseq_l6Dec04:*

- 1: Geneseqp1980s:*
- 2: Geneseqp1980s:*
- 3: Geneseqp2000s:*
- 4: Geneseqp2001s:*
- 5: Geneseqp2002s:*
- 6: Geneseqp2003as:*
- 7: Geneseqp2003bs:*
- 8: Geneseqp2004s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	2080	100.0	383	2	AAY08660 WO9927094
2	2080	100.0	383	2	AAY08657 Human tra
3	2080	100.0	383	2	AAY13390 Amino aci
4	2080	100.0	383	3	AAY88277 Human TAN
5	2080	100.0	383	3	AAY87270 Human sig
6	2080	100.0	383	3	AAY53627 A bone ma
7	2080	100.0	383	3	AAB25618 Protein e
8	2080	100.0	383	3	AAB25592 Protein e
9	2080	100.0	383	3	ADC78573 Human PRO
10	2080	100.0	383	4	AAB80258 Human PRO
11	2080	100.0	383	4	AAB48974 Human Zai
12	2080	100.0	383	4	AAB29048 Human PRO
13	2080	100.0	383	6	ABU58424 Human PRO
14	2080	100.0	383	6	ABU71636 Human PRO
15	2080	100.0	383	6	ABU87972 Novel hum
16	2080	100.0	383	6	ABU84287 Human sec
17	2080	100.0	383	6	ABR66161 Human sec
18	2080	100.0	383	6	ABR65551 Human sec
19	2080	100.0	383	6	ABU99491 Human sec
20	2080	100.0	383	6	ABU82730 Human PRO
21	2080	100.0	383	6	ABU89851 Novel hum
22	2080	100.0	383	6	ABU71491 Human PRO
23	2080	100.0	383	6	ABR68100 Human sec
24	2080	100.0	383	6	ABU96153 Novel hum
25	2080	100.0	383	6	ABU92584 Human sec

26	2080	100.0	383	6	ABO08661	AbO08661 Human sec
27	2080	100.0	383	6	ABO02713	AbO02713 Human sec
28	2080	100.0	383	6	ABR74867	ABr74867 Human sec
29	2080	100.0	383	6	ABR94629	ABr94629 Human sec
30	2080	100.0	383	6	ABU85602	ABu85602 Human PRO
31	2080	100.0	383	6	ABU98762	ABu98762 Novel hum
32	2080	100.0	383	6	ABU97977	ABu97977 Novel hum
33	2080	100.0	383	6	ABU91683	ABu91683 Novel hum
34	2080	100.0	383	6	ABU71937	ABu71937 Human sec
35	2080	100.0	383	6	ABU89376	ABu89376 Human PRO
36	2080	100.0	383	6	ABU86217	ABu86217 Human sec
37	2080	100.0	383	6	ABU67430	ABu67430 Human sec
38	2080	100.0	383	6	ABU80458	ABu80458 Human PRO
39	2080	100.0	383	6	ABO01820	ABO01820 Novel hum
40	2080	100.0	383	6	ABR99376	ABr99376 Human sec
41	2080	100.0	383	6	ABR98766	ABr98766 Human sec
42	2080	100.0	383	6	ABO16289	ABO16289 Human sec
43	2080	100.0	383	6	ABR92189	ABr92189 Human sec
44	2080	100.0	383	6	ABO18830	ABO18830 Human sec
45	2080	100.0	383	6	ABR78251	ABr78251 Human sec

ALIGNMENTS

RESULT 1
AAY08660
ID AAY08660 standard; protein; 383 AA.

XX AAY08660;

DT 09-AUG-1999 (first entry)

XX WO9927094 Seq ID 12.

XX Transmembrane domain; human; nutrition; cytokine; cell differentiation;
KW immune stimulation; immune suppression; haematopoiesis; activin;
KW regulatory tissue growth; inhibin; chemostatic; chemokinetic;
KW haemostatic; thrombolytic; tumour inhibitor; anti-inflammatory;
KW gene therapy; screening.

XX Homo sapiens.

XX WO9927094-A2.

XX 03-JUN-1999.

XX 20-NOV-1998; 98WO-JP005238.

XX 25-NOV-1997; 97JP-00323129.

XX (SAGA) SAGAMI CHEM RES CENT.

PA (PROT-) PROTEGENE INC.

XX Kato S, Kimura T, Sekine S;

PI WPI; 1999-357835/30.

XX Novel proteins containing transmembrane domains, useful as anti-

DR inflammatories, immune stimulators/suppressors and tissue growth

PT compounds.

XX Disclosure; Page 87-89; 89pp; English.

XX This invention describes novel human transmembrane containing proteins

CC and their encoding nucleic acids. Although no specific use is given for

CC the proteins, they may have a range of activities selected from

CC nutritional uses, cytokine and cell differentiation, immune

CC stimulation/suppression, haematopoiesis regulatory, tissue growth,

CC activin/inhibin, chemostatic/chemokinetic, haemostatic/thrombolytic,

CC receptor/ligand, tumour inhibitor, anti-inflammatory and other undefined

CC activities. The cDNAs can be utilized as probes for gene diagnosis and as

CC gene sources for gene therapy. The cDNAs can also be used for large scale

CC expression of proteins. The transformed cells can be used for detection
CC of the corresponding ligands and for screening of novel low-molecular
CC pharmaceuticals
XX
SQ Sequence 383 AA;

Query Match 100.0%; Score 2080; DB 2; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.8e-145;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGQCHKGTPPTYEEAKQYLSYETLYANGSRSTETQVGIYILSSSGDGAQHRDGS 120
DB 61 VSSSCGQCHKGTPPTYEEAKQYLSYETLYANGSRSTETQVGIYILSSSGDGAQHRDGS 120

QY 121 SGKRRKRQIYGDSRFSIFGKDFLLNYPFSTSVKLSGTGCTGLVAEKHVLTAACHIDG 180
DB 121 SGKRRKRQIYGDSRFSIFGKDFLLNYPFSTSVKLSGTGCTGLVAEKHVLTAACHIDG 180

QY 181 KTYVKGTKLRVGLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240
DB 181 KTYVKGTKLRVGLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240

QY 241 IGMDDYALLELKKPKRKFPMKIGVSPPAKQLPGRIHFSGYNDNRPGNLVYRFGDVKDE 300
DB 241 IGMDDYALLELKKPKRKFPMKIGVSPPAKQLPGRIHFSGYNDNRPGNLVYRFGDVKDE 300

QY 301 TYDLLYQCCDAQPGASGSGVYVRMWKQKQKWERKIIGIFSGHWDMDNGSPQDFNVAVR 360
DB 301 TYDLLYQCCDAQPGASGSGVYVRMWKQKQKWERKIIGIFSGHWDMDNGSPQDFNVAVR 360

QY 361 ITPLKYAQCWIKGNLYDCREG 383
DB 361 ITPLKYAQCWIKGNLYDCREG 383

RESULT 2
AAY08657
ID AAY08657 standard; protein; 383 AA.
AC AAY08657;
XX
DT 09-AUG-1999 (first entry)
XX
DE Human transmembrane domain containing protein from clone HF10493.
XX
KW Transmembrane domain; human; nutrition; cytokine; cell differentiation;
KW immune stimulation; immune suppression; haematopoiesis; activin;
KW regulatory tissue growth; inhibin; chemostatic; chemokinetic;
KW haemostatic; thrombolytic; tumour inhibitor; anti-inflammatory;
KW gene therapy; screening.
XX
OS Homo sapiens.
XX
PN WO9927094-A2.
XX
PD 03-JUN-1999.
XX
PF 20-NOV-1998; 98WO-JP005238.
XX
PR 25-NOV-1997; 97JP-00323129.
XX
PA (SAGA) SAGAMI CHEM RES CENT.
PA (PROT-) PROTEGENE INC.
XX
PI Kato S, Kimura T, Sekine S;
XX
DR MPI; 1999-357835/30.
DR N-PSDB; AAX77690, AAX77691.
XX

PT Novel proteins containing transmembrane domains, useful as anti-inflammatory, immune stimulators/suppressors and tissue growth compounds.
PT
XX
PS Claim 1; Page 68-69; 89pp; English.
XX
CC This invention describes novel human transmembrane containing proteins and their encoding nucleic acids. Although no specific use is given for the proteins, they may have a range of activities selected from CC nutritional uses, cytokine and cell differentiation, immune CC stimulation/suppression, haematopoiesis regulatory, tissue growth, CC activin/inhibin, chemostatic/chemokinetic, haemostatic/thrombolytic, CC receptor/ligand, tumour inhibitor, anti-inflammatory and other undefined CC activities. The cDNAs can be utilized as probes for gene diagnosis and as CC gene sources for gene therapy. The cDNAs can also be used for large scale CC expression of proteins. The transformed cells can be used for detection CC of the corresponding ligands and for screening of novel low-molecular CC pharmaceuticals
XX
SQ Sequence 383 AA;

Query Match 100.0%; Score 2080; DB 2; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.8e-145;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGQCHKGTPPTYEEAKQYLSYETLYANGSRSTETQVGIYILSSSGDGAQHRDGS 120
DB 61 VSSSCGQCHKGTPPTYEEAKQYLSYETLYANGSRSTETQVGIYILSSSGDGAQHRDGS 120

QY 121 SGKRRKRQIYGDSRFSIFGKDFLLNYPFSTSVKLSGTGCTGLVAEKHVLTAACHIDG 180
DB 121 SGKRRKRQIYGDSRFSIFGKDFLLNYPFSTSVKLSGTGCTGLVAEKHVLTAACHIDG 180

QY 181 KTYVKGTKLRVGLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240
DB 181 KTYVKGTKLRVGLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240

QY 241 IGMDDYALLELKKPKRKFPMKIGVSPPAKQLPGRIHFSGYNDNRPGNLVYRFGDVKDE 300
DB 241 IGMDDYALLELKKPKRKFPMKIGVSPPAKQLPGRIHFSGYNDNRPGNLVYRFGDVKDE 300

QY 301 TYDLLYQCCDAQPGASGSGVYVRMWKQKQKWERKIIGIFSGHWDMDNGSPQDFNVAVR 360
DB 301 TYDLLYQCCDAQPGASGSGVYVRMWKQKQKWERKIIGIFSGHWDMDNGSPQDFNVAVR 360

QY 361 ITPLKYAQCWIKGNLYDCREG 383
DB 361 ITPLKYAQCWIKGNLYDCREG 383

RESULT 3
AAY13390
ID AAY13390 standard; protein; 383 AA.
XX
AC AAY13390;
XX
DT 25-JUN-1999 (first entry)
XX
DE Amino acid sequence of protein PRO307.
XX
KW Secreted protein; transmembrane protein; human; enterocolitis;
KW Zollinger-Ellison syndrome; gastrointestinal ulceration;
KW congenital microvillus atrophy; skin disease; cell growth;
KW abnormal keratinocyte differentiation; psoriasis; epithelial cancer;
KW Parkinson's disease; Alzheimer's disease; ALS; neuropathy; fibromodulin;
KW dermal scarring; Usher Syndrome; Atrophia areata; anti-thrombotic;
KW wound healing; tissue repair.
XX
OS Homo sapiens.

XX PF 30-SEP-1999; 99WO-US022817.
 XX XX WPI: 2000-293144/25
 PR PR N-PSDB; AAA39945, AAA39946.
 XX 02-OCT-1998; 98US-00164220.
 XX 02-OCT-1998; 98US-00164169.
 PA (MILL-) MILLENNIUM BIOTHERAPEUTICS INC.
 XX Barnes TM;
 PI WPI: 2000-293144/25
 DR N-PSDB; AAA39945, AAA39946.
 DR
 XX Isolated nucleic acids encoding TANGO polypeptides useful for preventing,
 PT diagnosing and treating diseases associated with inappropriate protein
 PT expression.
 XX
 PS Claim 9; Fig 13; 249pp; English.
 XX
 XX This invention describes novel human and murine nucleic acids encoding
 CC TANGO polypeptides (which are either wholly secreted or transmembrane
 CC proteins) which can be used for gene therapy and/or vaccination. The
 CC peptides are designated TANGO 180 to TANGO 189 and TANGO 215. The nucleic
 CC acids may be used to produce TANGO 180 to TANGO 189 and TANGO 215
 CC polypeptides according to standard recombinant DNA methodologies. They
 CC may also be used to detect and quantify the presence of TANGO nucleic
 CC acids in a sample and therefore identify or diagnose diseases associated
 CC with inappropriate TANGO expression (e.g. diseases related to over or
 CC under expression of the polypeptides or the expression of inactive
 CC polypeptides). The nucleic acids and the polypeptides they encode may be
 CC used according to standard gene therapy protocols, to treat diseases
 CC associated with inappropriate TANGO expression by supplementing a
 CC patient's own production of the polypeptide of to rectify mutations that
 CC may result in expression of an abnormally active polypeptide. The
 CC polypeptides may also be used to identify and produce agonists and
 CC antagonists of TANGO expression and activity which may be used to
 CC modulate TANGO related processes and diseases. The polypeptides are
 CC particularly useful for use as antigens for producing antibodies to TANGO
 CC proteins which may be used for inhibiting the activity of TANGO proteins.
 CC They may also be used to detect and quantify the presence of TANGO
 CC proteins in samples and therefore identify patients in whom the protein
 CC is over- or under-expressed. This sequence represents the human TANGO 186
 CC protein described in the method of the invention
 XX
 SQ Sequence 383 AA;
 Query Match 100.0%; Score 2080; DB 3; Length 383;
 Best Local Similarity 100.0%; Pred. No. 1.8e-145;
 Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Qy 1 MAGIPGLLFLFLLCVAGQVSPYSAPWKPETWPAYRLPVVLPOSTLNLAKPFGAEAKLE 60
 Db 1 MAGIPGLLFLFLLCVAGQVSPYSAPWKPETWPAYRLPVVLPOSTLNLAKPFGAEAKLE 60
 Qy 61 VSSCGPQCHKGTPLPYERAKQVLSYETLYANGSRTEQVGIYILSSSGDGAQRDSDGS 120
 Db 61 VSSCGPQCHKGTPLPYERAKQVLSYETLYANGSRTEQVGIYILSSSGDGAQRDSDGS 120
 Qy 121 SGKSRKROLYGVDSRESIFGKDFLLNYPSTSVKLSGTGCTGLVAEKHVLTAACHIDHG 180
 Db 121 SGKSRKROLYGVDSRESIFGKDFLLNYPSTSVKLSGTGCTGLVAEKHVLTAACHIDHG 180
 Qy 181 KTVYKGTQKLVRGPKFKFDGGRGANDSTSAMPEQMKFQWIRKRVTHVPKGMKGNAND 240
 Db 181 KTVYKGTQKLVRGPKFKFDGGRGANDSTSAMPEQMKFQWIRKRVTHVPKGMKGNAND 240
 Qy 241 IGMDDYVALLLELKKPHRKPKMKIGVSPAPKQLPGRRIHFGSYNDNRPNLVYRFDVWDE 300
 Db 241 IGMDDYVALLLELKKPHRKPKMKIGVSPAPKQLPGRRIHFGSYNDNRPNLVYRFDVWDE 300
 Qy 301 TYDLLYQCCDAQPCGASGVYVRWKKQKQKWKRIIGFSGHWDVMDNGSPQDFNVAVR 360
 Db 301 TYDLLYQCCDAQPCGASGVYVRWKKQKQKWKRIIGFSGHWDVMDNGSPQDFNVAVR 360

Qy 361 ITPLKYAQICYWIKGNYLDCREG 383
 |||||
 Db 361 ITPLKYAQICYWIKGNYLDCREG 383
 |||||
 RESULT 5
 AAY87270
 ID AAY87270 standard; protein; 383 AA.
 XX
 AC AAY87270;
 XX
 DT 11-MAY-2000 (first entry)
 XX
 DE Human signal peptide containing protein HSPP-47 SEQ ID NO:47.
 XX
 KW Human; signal peptide-containing protein; HSPP; diagnosis; cancer;
 KW inflammation; cardiovascular disease; anticancer; anti-inflammatory;
 KW antimicrobial; nontropic; neuroprotective; cardiovascular; hepatotropic;
 KW antiasthmatic; gene therapy; cell proliferation; neurological disorder;
 KW reproductive disorder; developmental disorder; arteriosclerosis;
 KW cirrhosis; psoriasis; acquired immune deficiency syndrome; anaemia;
 KW asthma; Crohn's disease; infection; Alzheimer's disease; schizophrenia;
 KW Parkinson's disease; Huntington's diseases; ovulatory defect;
 muscular dystrophy.
 OS Homo sapiens.
 PN WO200000610-A2.
 XX
 PD 06-JAN-2000.
 XX
 PF 25-JUN-1999; 99WO-US014484.
 XX
 PR 26-JUN-1998; 98US-0090762P.
 PR 31-JUL-1998; 98US-0094983P.
 PR 01-OCT-1998; 98US-0102886P.
 PR 11-DEC-1998; 98US-0112129P.
 XX
 PA (INCY-) INCYTE PHARM INC.
 XX
 PI Lal P, Tang YT, Gorgone GA, Corley NC, Guegler KJ, Baughn MR;
 PI Akerblom IE, Au-Young J, Yue H, Patterson C, Reddy R, Hillman JL;
 PI Bandman O;
 XX
 DR WPI: 2000-160673/14.
 DR N-PSDB; AA298155.
 XX
 PT New human signal peptide-containing proteins useful in treatment,
 PT prevention and diagnosis of e.g. cancer, inflammation and cardiovascular
 PT disease.
 XX
 PS Claim 1; Page 192-193; 327pp; English.
 XX
 CC AA298109 to AA298242 encode AAY87224 to AAY87357 which represent the
 CC human signal peptide-containing proteins HSPP-1 to HSPP-134. HSPPs have
 CC anticancer, anti-inflammatory, antimicrobial, nontropic, hepatotropic,
 CC neuroprotective, cardiovascular and antiasthmatic activities, and can be
 CC used in gene therapy. HSPPs can be used to treat or prevent disorders
 CC associated with decreased activity or function of HSPP. Antagonists of
 CC HSPP are used to treat or prevent disorders associated with increased
 CC activity or function of HSPP. Such diseases include cell proliferation
 CC (including cancer), inflammation, cardiovascular, neurological,
 CC reproductive or developmental disorders, (e.g. arteriosclerosis,
 CC cirrhosis, psoriasis, acquired immune deficiency syndrome, congestive or
 CC ischaemic heart disease, Alzheimer's, Parkinson's or Huntington's
 CC diseases, schizophrenia, ovulatory defects, muscular dystrophy). HSPP
 CC nucleic acids can be used for the recombinant production of HSPP, for
 CC detecting HSPP in standard hybridisation and amplification assays (for
 CC diagnosis and monitoring), in gene therapy, as antisense, triplex-forming
 CC or ribozyme therapeutics, for detecting related sequences or genetic
 CC variations, and for chromosomal mapping. HSPP are also used to raise

CC specific antibodies (Ab) and to screen for agonists and antagonists
CC (potential therapeutic agents). Ab are used to diagnose, or monitor, HSPP
CC -related diseases (in usual immunoassays), as therapeutic antagonists, in
CC competitive drug screens, and for purification of HSPP from natural
CC sources
XX
XX Sequence 383 AA;
SQ
Query Match 100.0%; Score 2080; DB 3; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.8e-145;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 MAGIPGLLFLFFLLCAVGVSPYSAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFFLLCAVGVSPYSAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
Qy 61 VSSSCGPQCHKGTPPTYBEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS 120
Db 61 VSSSCGPQCHKGTPPTYBEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS 120
Qy 121 SGKSRKRQIYGDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Qy 181 KTYVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVTRTHVPKGIKGNAND 240
Db 181 KTYVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVTRTHVPKGIKGNAND 240
Qy 241 IGMDDYDYLLELKKPHKRFKMGIVSPPAKQLPGGRIFHSFYNDNDRPGLNLYRFDCKDE 300
Db 241 IGMDDYDYLLELKKPHKRFKMGIVSPPAKQLPGGRIFHSFYNDNDRPGLNLYRFDCKDE 300
Qy 301 TYDLYQQCDAQPGASGSGVYVVMWKRQOQKWERKIIGIFSGHQMNGSPQDFNVAVR 360
Db 301 TYDLYQQCDAQPGASGSGVYVVMWKRQOQKWERKIIGIFSGHQMNGSPQDFNVAVR 360
Qy 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383
RESULT 6
AAV53627
ID AAY53627 standard; protein; 383 AA.
XX
AC AAY53627;
XX
DT 22-FEB-2000 (first entry)
XX
DE A bone marrow secreted protein designated BMS192.
XX
KW Bone marrow secreted protein; bone marrow stromal cell; cytokine;
KW cell proliferation; cell differentiation; hematopoiesis; anaemia;
KW myeloid cell deficiency; lymphoid cell deficiency; myeloid cell;
KW erythroid progenitor cell; colony stimulating factor; granulocyte;
KW monocyte; macrophage; myelo-suppression; megakaryocyte; platelet;
KW platelet disorder; thrombocytopenia; hematopoietic stem cell;
KW stem cell disorder; aplastic anaemia; bone differentiation;
KW paroxysmal nocturnal hemoglobinuria; bone growth; cartilage; tendon;
KW ligament; nerve; wound healing; tissue repair; burn; incision; ulcer;
KW bone fracture; cartilage damage; artificial joint.
XX
OS Homo sapiens.
XX
FH Key Location/Qualifiers
FT Peptide 1..19
FT /note= "signal peptide"
XX
XX W09933979-A2.
XX
XX 08-JUL-1999.
XX
XX 18-DEC-1998; 98WO-US027008.

XX 30-DEC-1997; 97US-0068958P.
PR 24-SEP-1998; 98US-0101603P.
PR 30-SEP-1998; 98US-0102540P.
XX
PA (CHIR) CHIRON CORP.
XX
XX Lin H, Cao L;
XX
XX WPI; 2000-038344/03.
DR N-PSDB; AA236233.
XX
XX New isolated human polynucleotide and secreted proteins can induce
XX production of other cytokines in certain cell populations.
PS Claim 2; Page 81-82; 120pp; English.
XX
XX AAY53622-43 represent bone marrow secreted proteins of human bone marrow
XX stromal cells. The proteins can exhibit cytokine, cell proliferation, or
XX cell differentiation activity (either inducing or inhibiting). They can
XX be used to support colony forming cells or factor-dependent cell lines,
XX to regulate hematopoiesis, and to treat myeloid or lymphoid cell
XX deficiencies. In addition, they may be used to support the growth and
XX proliferation of erythroid progenitor cells, and to treat various
XX anaemias. They can have colony stimulating factor (CSF) activity and can
XX be used to support the growth and proliferation of myeloid cells such as
XX granulocytes, monocytes or macrophages, to prevent or treat myelo-
XX suppression, to support the growth and proliferation of megakaryocytes
XX and platelets, thereby allowing prevention or treatment of platelet
XX disorders such as thrombocytopenia, to support the growth and
XX proliferation of hematopoietic stem cells, either in place of or in
XX conjunction with platelet transfusions, to treat stem cell disorders,
XX such as aplastic anaemia and paroxysmal nocturnal hemoglobinuria, or to
XX repopulate the stem cell compartment after irradiation or chemotherapy.
XX They can be used for growth or differentiation of bone, cartilage,
XX tendon, ligament, or nerve tissue, as well as for wound healing and
XX tissue repair and replacement, and in the treatment of burns, incisions
XX and ulcers; to induce cartilage and/or bone growth in circumstances where
XX bone is not normally formed and thus have an application in healing bone
XX fractures and cartilage damage or defects, prophylactic use in fracture
XX reduction and also in the improved fixation of artificial joints
XX
SQ Sequence 383 AA;
Query Match 100.0%; Score 2080; DB 3; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.8e-145;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 MAGIPGLLFLFFLLCAVGVSPYSAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFFLLCAVGVSPYSAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
Qy 61 VSSSCGPQCHKGTPPTYBEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS 120
Db 61 VSSSCGPQCHKGTPPTYBEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS 120
Qy 121 SGKSRKRQIYGDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Qy 181 KTYVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVTRTHVPKGIKGNAND 240
Db 181 KTYVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVTRTHVPKGIKGNAND 240
Qy 241 IGMDDYDYLLELKKPHKRFKMGIVSPPAKQLPGGRIFHSFYNDNDRPGLNLYRFDCKDE 300
Db 241 IGMDDYDYLLELKKPHKRFKMGIVSPPAKQLPGGRIFHSFYNDNDRPGLNLYRFDCKDE 300
Qy 301 TYDLYQQCDAQPGASGSGVYVVMWKRQOQKWERKIIGIFSGHQMNGSPQDFNVAVR 360
Db 301 TYDLYQQCDAQPGASGSGVYVVMWKRQOQKWERKIIGIFSGHQMNGSPQDFNVAVR 360
Qy 361 ITPLKYAQICYWIKGNLYDCREG 383

Db	361	ITPLKYAQICYWIKGNLYDCREG 383	Query Match 100.0%; Score 2080; DB 3; Length 383; Best Local Similarity 100.0%; Pred. No. 1.8e-145; Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
RESULT 7			
AAAB25518			
ID	AAAB25518	standard; protein; 383 AA.	
XX	AC	AAAB25518;	
XX	AC	AAAB25518;	
XX	DT	21-NOV-2000 (first entry)	
XX	DT	21-NOV-2000 (first entry)	
XX	DE	Protein encoded by human secreted protein gene #10.	
XX	XX	Secreted protein; immunosuppressant; anti-inflammatory; antiarthritic;	
XX	KW	antirheumatic; dermatological; antiproliferative; antiarteriosclerotic;	
XX	KW	anticancer; vulnery; antiviral; antibacterial; antifungal;	
XX	KW	immune disorder; Addison's disease; rheumatoid arthritis; dermatitis;	
XX	KW	multiple sclerosis; inflammatory disorder; inflammatory bowel disease;	
XX	KW	Crohn's disease; nephritis; hyperproliferative disorder;	
XX	KW	cardiovascular disorder; coronary arteriosclerosis; myocarditis; cancer;	
XX	KW	melanoma; lymphoma; wound healing; human; chromosome 12.	
XX	OS	Homo sapiens.	
XX	XX	WO200029435-A1.	
XX	PN	25-MAY-2000.	
XX	PD	25-MAY-2000.	
XX	XX	27-OCT-1999; 99WO-US025031.	
XX	PF	27-OCT-1999; 99WO-US025031.	
XX	XX	28-OCT-1998; 98US-0105971P.	
XX	PR	28-OCT-1998; 98US-0105971P.	
XX	XX	(HUMA-) HUMAN GENOME SCI INC.	
XX	PA	(HUMA-) HUMAN GENOME SCI INC.	
XX	PI	Ni J, Ruben SM, Olsen HS, Young PE, Kenny JJ, Moore PA, Wei Y;	
XX	PI	Greene JM;	
XX	XX	WPI; 2000-387742/33.	
XX	DR	Isolated nucleic acid molecules encoding human secreted proteins are used	
XX	XX	for the prevention, amelioration and treatment of autoimmune, cancer,	
XX	PT	inflammatory, hyperproliferative and cardiovascular disorders, cancer,	
XX	PT	wounds, and infectious diseases.	
XX	PS	Disclosure; Page 169; 803pp; English.	
XX	XX	The present invention relates to 12 secreted human proteins and the	
XX	CC	nucleotide sequences encoding them. The polynucleotide sequences given in	
XX	CC	AAAB0606-A80623 encode the 12 secreted protein sequences given in	
XX	CC	AAAB25576-B25593. The human secreted proteins have various activities	
XX	CC	dependent on the tissues in which they are expressed. Examples of the	
XX	CC	activities of the proteins include: immunosuppressant; anti-inflammatory;	
XX	CC	antiarthritic; antirheumatic; dermatological; antiproliferative;	
XX	CC	antiarteriosclerotic; anticancer; vulnery; antiviral; antibacterial;	
XX	CC	and antifungal activity. The proteins, polypeptides, agonists and	
XX	CC	antagonists may be used to treat prevent and/or diagnose various disease,	
XX	CC	disorders and conditions examples of which include: immune disorders e.g.	
XX	CC	Addison's disease, rheumatoid arthritis, dermatitis, and multiple	
XX	CC	sclerosis; inflammatory disorders e.g. inflammatory bowel disease,	
XX	CC	Crohn's disease and nephritis; hyperproliferative disorders such as	
XX	CC	paraproteinemia and purpura; cardiovascular disorders e.g. coronary	
XX	CC	arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The	
XX	CC	proteins and polynucleotide sequences may also be used in wound healing	
XX	CC	and the treatment of infectious diseases. The human secreted protein gene	
XX	CC	#10 and protein sequences are represented in sequences AAAB0615 and	
XX	CC	AAAB25585. Secreted protein gene #10 is located on chromosome 12.	
XX	CC	Sequences AAAB25516-B25518 represent alternative secreted protein #10	
XX	CC	sequences and AAAB0669-A80676 represent genes which are related to the	
XX	CC	secreted protein gene#10	
XX	XX	Sequence 383 AA;	

QY	1	MAGIPGLLLFLPFLICAVGVSPYSAPWKPPTWPAAYRLPVVLPSTLNLAKEPFGAEAKLE 60	
Db	1	MAGIPGLLLFLPFLICAVGVSPYSAPWKPPTWPAAYRLPVVLPSTLNLAKEPFGAEAKLE 60	
QY	61	VSSSCGPGQCHKGTPPTLYBEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDGS 120	
Db	61	VSSSCGPGQCHKGTPPTLYBEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDGS 120	
QY	121	SGKSRKROIYGYDSRFSIFGKDFILNYPSTSVKLSCTGTLVAEKHVLTAACHIDG 180	
Db	121	SGKSRKROIYGYDSRFSIFGKDFILNYPSTSVKLSCTGTLVAEKHVLTAACHIDG 180	
QY	181	KTVVKTQKLRVGFLLPKPKFGDGGGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240	
Db	181	KTVVKTQKLRVGFLLPKPKFGDGGGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240	
QY	241	IGMDYDVALLELKKPHKRFMKIGVSPPAKQLPGRHFSGYDNDPRPNLVYRFDVKDE 300	
Db	241	IGMDYDVALLELKKPHKRFMKIGVSPPAKQLPGRHFSGYDNDPRPNLVYRFDVKDE 300	
QY	301	TYDLLYQCCDAQPGASGSGYVVRMWKQKQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360	
Db	301	TYDLLYQCCDAQPGASGSGYVVRMWKQKQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360	
QY	361	ITPLKYAQICYWIKGNLYDCREG 383	
Db	361	ITPLKYAQICYWIKGNLYDCREG 383	
RESULT 8			
AAAB25592			
ID	AAAB25592	standard; protein; 383 AA.	
XX	AC	AAAB25592;	
XX	DT	21-NOV-2000 (first entry)	
XX	DT	21-NOV-2000 (first entry)	
XX	DE	Protein encoded by human secreted protein gene #10 clone HUSQ05.	
XX	XX	Secreted protein; immunosuppressant; anti-inflammatory; antiarthritic;	
XX	KW	antirheumatic; dermatological; antiproliferative; antiarteriosclerotic;	
XX	KW	anticancer; vulnery; antiviral; antibacterial; antifungal;	
XX	KW	immune disorder; Addison's disease; rheumatoid arthritis; dermatitis;	
XX	KW	multiple sclerosis; inflammatory disorder; inflammatory bowel disease;	
XX	KW	Crohn's disease; nephritis; hyperproliferative disorder;	
XX	KW	cardiovascular disorder; coronary arteriosclerosis; myocarditis; cancer;	
XX	KW	melanoma; lymphoma; wound healing; human; chromosome 12.	
XX	OS	Homo sapiens.	
XX	XX	WO200029435-A1.	
XX	PN	25-MAY-2000.	
XX	PD	25-MAY-2000.	
XX	XX	27-OCT-1999; 99WO-US025031.	
XX	PF	27-OCT-1999; 99WO-US025031.	
XX	XX	28-OCT-1998; 98US-0105971P.	
XX	PR	28-OCT-1998; 98US-0105971P.	
XX	XX	(HUMA-) HUMAN GENOME SCI INC.	
XX	PA	(HUMA-) HUMAN GENOME SCI INC.	
XX	PI	Ni J, Ruben SM, Olsen HS, Young PE, Kenny JJ, Moore PA, Wei Y;	
XX	PI	Greene JM;	
XX	XX	WPI; 2000-387742/33.	
XX	DR	Isolated nucleic acid molecules encoding human secreted proteins are used	
XX	XX	for the prevention, amelioration and treatment of autoimmune, cancer,	
XX	PT	inflammatory, hyperproliferative and cardiovascular disorders, cancer,	
XX	PT	wounds, and infectious diseases.	

XX Claim 1; Page 684-685; 803pp; English.

XX The present invention relates to 12 secreted human proteins and the

CC nucleotide sequences encoding them. The polynucleotide sequences given in

CC AAB0606-A80623 encode the 12 secreted protein sequences given in

CC AAB25576-B25593. The human secreted proteins have various activities

CC dependent on the tissues in which they are expressed. Examples of the

CC activities of the proteins include: immunosuppressant; anti-inflammatory;

CC antiarthritic; antirheumatic; dermatological; antiproliferative;

CC antiarteriosclerotic; anticancer; vulnerary; antiviral; antibacterial;

CC and antifungal activity. The proteins, polypeptides, agonists and

CC antagonists may be used to treat prevent and/or diagnose various disease,

CC disorders and conditions examples of which include: immune disorders e.g.

CC Addison's disease, rheumatoid arthritis, dermatitis, and multiple

CC sclerosis; inflammatory disorders e.g. inflammatory bowel disease,

CC Crohn's disease and nephritis; hyperproliferative disorders such as

CC paraproteinemia and purpura; cardiovascular disorders e.g. coronary

CC arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The

CC proteins and polynucleotide sequences may also be used in wound healing

CC and the treatment of infectious diseases. The human secreted protein gene

CC #10 and protein sequences are represented in sequences AAB0606 and

CC AAB25585. Secreted protein gene #10 is located on chromosome 12.

CC Sequences AAB25616-B25618 represent alternative secreted protein #10

CC sequences and AAB0669-A80676 represent genes which are related to the

CC secreted protein gene#10

XX Sequence 383 AA;

Query Match 100.0%; Score 2080; DB 3; Length 383;

Best Local Similarity 100.0%; Pred. No. 1.8e-145;

Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLLFLLLCAVGVSPYSAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60

DB 1 MAGIPGLLLFLLLCAVGVSPYSAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGQCHKGTPPLTYEAKQVLSYETLYANGSRRTQVGIYILSSSGDGAQHRDGS 120

DB 61 VSSSCGQCHKGTPPLTYEAKQVLSYETLYANGSRRTQVGIYILSSSGDGAQHRDGS 120

QY 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLTGCTGLVAEKHVLTAACHIDG 180

DB 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLTGCTGLVAEKHVLTAACHIDG 180

QY 181 KTVYKGTQKLRVGLPKPKFGKGGRGANDSTSAMPEQMKFQWIRKTRTHVPKGIKGNAND 240

DB 181 KTVYKGTQKLRVGLPKPKFGKGGRGANDSTSAMPEQMKFQWIRKTRTHVPKGIKGNAND 240

QY 241 IGMDYDVALLELKKPKHKKFKMKGIVSPPAKQLPGGRIFHSGYDNDPGLNLYRFDVKDE 300

DB 241 IGMDYDVALLELKKPKHKKFKMKGIVSPPAKQLPGGRIFHSGYDNDPGLNLYRFDVKDE 300

QY 301 TYDLYQQCDAQPCASGSGVYVWRKQKQKWERKIIGFSGHGWDMNGSPQDFNVAVR 360

DB 301 TYDLYQQCDAQPCASGSGVYVWRKQKQKWERKIIGFSGHGWDMNGSPQDFNVAVR 360

QY 361 ITPLKYAICWIKGNLYDCREG 383

DB 361 ITPLKYAICWIKGNLYDCREG 383

RESULT 9

ADC78573

ID ADC78573

XX standard; protein; 383 AA.

AC ADC78573;

XX

01-JAN-2004 (first entry)

DT Human PRO307 protein.

DE

XX antiinflammatory; antiulcer; cytostatic; antipsoriatic; antiparkinsonian;

KW neurotropic; neuroprotective; vasotropic; chemotactic; angiogenic;

KW neurotrophic; osteopathic; antiasthmatic; antiarthritic; antirheumatic;

KW antiarteriosclerotic; cardiant; antidiabetic; cerebroprotective;

KW thrombolytic; immunomodulator; enterocolitis; Zollinger-Ellison syndrome;

KW gastrointestinal ulceration; psoriasis; cancer; Parkinson's disease;

KW Alzheimer's; ALS; neuropathy; dermal scarring; wound healing;

KW nerve repair; thrombosis; bone; cartilage formation; angiogenesis;

KW asthma; rheumatoid arthritis; multiple sclerosis; inflammatory disorder;

KW atherosclerosis; cardiac injury; infertility; premature aging; AIDS;

KW diabetes; stroke; gene therapy; transgenic; PRO; human.

OS Homo sapiens.

XX WO200015796-A2.

PN 23-MAR-2000.

XX 15-SEP-1999; 99WO-US021090.

PR 16-SEP-1998; 98WO-US019330.

XX (GETH) GENENTECH INC.

XX Chen J, Goddard A, Gurney AL, Hillan K, Pennica D, Wood WJ;

PI Yuan J;

XX WPI; 2000-271434/23.

DR N-PSDB; ADC78572.

XX Novel nucleic acids encoding secreted and transmembrane polypeptides with

PT homology, e.g. to growth and cancer-associated antigens.

PS Claim 12; SEQ ID NO 261; 355pp; English.

XX The invention relates to a novel nucleic acid encoding a PRO polypeptide.

CC The polypeptides and polynucleotides of the invention may be useful as

CC research tools and as therapeutics for treating enterocolitis, Zollinger-

CC Ellison syndrome, gastrointestinal ulceration, psoriasis, cancer,

CC Parkinson's disease, Alzheimer's disease, ALS, neuropathies, dermal

CC scarring and wound healing, nerve repair, thrombosis, bone and/or

CC cartilage formation, angiogenesis, asthma, rheumatoid arthritis, multiple

CC sclerosis, inflammatory disorders, atherosclerosis, cardiac injury,

CC infertility, premature aging, AIDS, diabetes complications and stroke.

CC The molecules may also be utilised during gene therapy procedures and

CC transgenic animal production. The current sequence is that of the human

XX PRO protein of the invention.

SQ Sequence 383 AA;

Query Match 100.0%; Score 2080; DB 3; Length 383;

Best Local Similarity 100.0%; Pred. No. 1.8e-145;

Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLLFLLLCAVGVSPYSAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60

DB 1 MAGIPGLLLFLLLCAVGVSPYSAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGQCHKGTPPLTYEAKQVLSYETLYANGSRRTQVGIYILSSSGDGAQHRDGS 120

DB 61 VSSSCGQCHKGTPPLTYEAKQVLSYETLYANGSRRTQVGIYILSSSGDGAQHRDGS 120

QY 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLTGCTGLVAEKHVLTAACHIDG 180

DB 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLTGCTGLVAEKHVLTAACHIDG 180

QY 181 KTVYKGTQKLRVGLPKPKFGKGGRGANDSTSAMPEQMKFQWIRKTRTHVPKGIKGNAND 240

DB 181 KTVYKGTQKLRVGLPKPKFGKGGRGANDSTSAMPEQMKFQWIRKTRTHVPKGIKGNAND 240

QY 241 IGMDYDVALLELKKPKHKKFKMKGIVSPPAKQLPGGRIFHSGYDNDPGLNLYRFDVKDE 300

DB 241 IGMDYDVALLELKKPKHKKFKMKGIVSPPAKQLPGGRIFHSGYDNDPGLNLYRFDVKDE 300

QY 301 TYDLLYQCDAPQASGSGVYVVMWKRQKWKRIIGIFSGHWDVMDNGSPQDFNVAVR 360
 Db TYDLLYQCDAPQASGSGVYVVMWKRQKWKRIIGIFSGHWDVMDNGSPQDFNVAVR 360
 QY 361 ITPLKYAQICWIKGNLYDCREG 383
 Db 361 ITPLKYAQICWIKGNLYDCREG 383

RESULT 10
 AAB80258
 ID AAB80258 standard; protein; 383 AA.
 XX
 AC AAB80258;
 XX
 DT 24-APR-2001 (first entry)
 XX
 DE Human PRO307 protein.
 XX
 KW Human; PRO; dermatological; antipsoriatic; cytostatic; antiinflammatory;
 KW antiparkinsonian nootropic; neuroprotective; vulnerary; cardiant;
 KW antiangiogenic; vasotropic; antiasthmatic; antirheumatic; cancer;
 KW antiarthritic; antinfertility; antidiabetic; antiviral; diabetes;
 KW ophthalmological; gene therapy; skin disease; gastrointestinal disorder;
 KW ischaemia; inflammation.
 OS
 XX Homo sapiens.
 XX
 PN WO200104311-A1.
 XX
 PD 18-JAN-2001.
 XX
 PF 22-FEB-2000; 2000WO-US004414.
 XX
 PR 07-JUL-1999; 99US-0143048P.
 PR 26-JUL-1999; 99US-0145698P.
 PR 28-JUL-1999; 99US-0146222P.
 PR 08-SEP-1999; 99WO-US020594.
 PR 13-SEP-1999; 99WO-US020944.
 PR 15-SEP-1999; 99WO-US021090.
 PR 15-SEP-1999; 99WO-US021547.
 PR 05-OCT-1999; 99WO-US023089.
 PR 29-NOV-1999; 99WO-US028214.
 PR 30-NOV-1999; 99WO-US028313.
 PR 02-DEC-1999; 99WO-US028564.
 PR 02-DEC-1999; 99WO-US028565.
 PR 16-DEC-1999; 99WO-US030095.
 PR 20-DEC-1999; 99WO-US030911.
 PR 20-DEC-1999; 99WO-US030999.
 PR 05-JAN-2000; 2000WO-US000219.
 XX
 PA (GETH) GENENTECH INC.
 XX
 PI Ashkenazi AJ, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
 PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
 PI Godowski PJ, Grimaldi CJ, Gurney AL, Hillan KJ, Kljavin IJ;
 PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
 PI Williams PM, Wood WI;
 XX
 DR WPI; 2001-081051/09.
 DR N-PSDB; AAF72419.
 XX
 PT Sixty one nucleic acids encoding PRO polypeptides which are useful in the
 PT treatment of skin diseases (e.g. psoriasis), cancers (e.g. lung squamous
 PT cell carcinoma) and neurodegenerative diseases (e.g. Alzheimer's
 PT disease).
 XX
 PS Claim 1; Fig 96; 393pp; English.
 XX
 CC The present sequence is one of sixty one novel secreted and transmembrane
 CC PRO polypeptides. The PRO polypeptides are useful for treating skin
 CC diseases (e.g. psoriasis), cancers (e.g. lung squamous cell carcinoma),
 CC gastrointestinal disorders (e.g. enterocolitis), neurodegenerative

CC diseases (e.g. Alzheimer's disease, Parkinson's disease), wound repair,
 CC cardiovascular disorders (e.g. endometrial bleeding angiogenesis,
 CC ischaemias such as coronary ischaemia, atherosclerosis), inflammatory
 CC disorders (e.g. asthma, rheumatoid arthritis, multiple sclerosis),
 CC infertility, AIDS and diabetes and retinal disorders such as retinitis
 CC pigmentosum. The PRO nucleic acids have applications in molecular
 CC biology, including use as hybridization probes, and in chromosome and
 CC gene mapping
 XX
 SQ Sequence 383 AA;
 Query Match 100.0%; Score 2080; DB 4; Length 383;
 Best Local Similarity 100.0%; Pred. No. 1.8e-145;
 Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MAGIPGLLLFLLPFLCAVGVSPYSAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
 Db 1 MAGIPGLLLFLLPFLCAVGVSPYSAPWKPWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
 QY 61 VSSSCGPQCHKGTPLPTYEAKQYLSYETLYANGSRRTETQVGIYILSSSGDGAQHRDGS 120
 Db 61 VSSSCGPQCHKGTPLPTYEAKQYLSYETLYANGSRRTETQVGIYILSSSGDGAQHRDGS 120
 QY 121 SGKSRRKROIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
 Db 121 SGKSRRKROIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
 QY 181 KTVVGTOKLRVGFLLPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
 Db 181 KTVVGTOKLRVGFLLPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
 QY 241 IGMDYDYLLELKKPHKRFKMGKIGVSPPAKQLPGGRIHFSGYDNDPRGNLVYRFDVKDE 300
 Db 241 IGMDYDYLLELKKPHKRFKMGKIGVSPPAKQLPGGRIHFSGYDNDPRGNLVYRFDVKDE 300
 QY 301 TYDLLYQCDAPQASGSGVYVVMWKRQKWKRIIGIFSGHWDVMDNGSPQDFNVAVR 360
 Db 301 TYDLLYQCDAPQASGSGVYVVMWKRQKWKRIIGIFSGHWDVMDNGSPQDFNVAVR 360
 QY 361 ITPLKYAQICWIKGNLYDCREG 383
 Db 361 ITPLKYAQICWIKGNLYDCREG 383

RESULT 11
 AAB48974
 ID AAB48974 standard; protein; 383 AA.
 XX
 AC AAB48974;
 XX
 DT 27-MAR-2001 (first entry)
 XX
 DE Human Zsig13 variant #3, SEQ ID NO:18.
 XX
 KW Human Zsig13; serine protease; chromosome 11q22.1; elastase homologue;
 KW glutamyl endopeptidase homologue; factor x homologue; trypsin homologue;
 KW trypsinogen homologue; mast cell protease homologue;
 KW collagenase homologue; protein degradation; food processing; brewing;
 KW alcohol production; laundry detergent component.
 XX
 OS Homo sapiens.
 XX
 PN US6153420-A.
 XX
 PD 28-NOV-2000.
 XX
 PF 04-MAY-1998; 98US-00072384.
 PR 24-APR-1997; 97US-0044185P.
 PR 17-APR-1998; 98US-00062142.
 XX
 PA (ZYMO) ZYMOGENETICS INC.
 XX

PI Sheppard PO;
 XX WPI; 2001-060090/07.
 DR N-PSDB; AAC91784.
 XX
 PT New isolated serine protease (designated Zsig13), useful in industrial
 PT processes to degrade unwanted proteins or alter the characteristics of
 PT protein-containing composition, as well as in industrial applications
 PT (e.g. brewing).
 XX
 XX Claim 1; Col 41-44; 26pp; English.
 XX
 CC The invention relates to human Zsig13 proteins (AAB48972-B48974), and to
 CC DNA encoding them (AAC91782-C91784). The invention also relates to
 CC expression vectors and host cells comprising a human Zsig13 DNA, and the
 CC recombinant production of a human Zsig13 protein or its precursor. Zsig13
 CC is a serine protease, and has significant homology to Bacillus
 CC licheniformis glutamyl endopeptidase, human clotting factor X, human
 CC elastase, rat mast cell protease, Streptomyces griseus trypsin, bovine
 CC trypsinogen, and Hypoderma lineatum collagenase. The gene encoding human
 CC Zsig13 is located on chromosome 11q22.1. Zsig13 is useful in industrial
 CC processes to degrade unwanted proteins or alter the characteristics of
 CC protein-containing compositions. It may also be used in industrial
 CC applications in which proteases are utilized, including food processing,
 CC brewing and alcohol production, and as a component of a laundry
 CC detergent. The present sequence represents a human Zsig13 variant
 XX
 XX Sequence 383 AA;
 SQ
 Query Match 100.0%; Score 2080; DB 4; Length 383;
 Best Local Similarity 100.0%; Pred. No. 1.8e-145;
 Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MAGIPGLLFLFLLCVAGQVSPYAPKPTPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
 Db 1 MAGIPGLLFLFLLCVAGQVSPYAPKPTPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
 QY 61 VSSSCGQCHGKGTPLTYEBAKQYLSYETLYANGSRRTETQVGIYILSSSGDGAQHRDGS 120
 Db 61 VSSSCGQCHGKGTPLTYEBAKQYLSYETLYANGSRRTETQVGIYILSSSGDGAQHRDGS 120
 QY 121 SGKRRKQIYGVDSRFSIFGKDFLLNYPSTSVKLTSTGCTGLVAEKHVLTAACHIDHG 180
 Db 121 SGKRRKQIYGVDSRFSIFGKDFLLNYPSTSVKLTSTGCTGLVAEKHVLTAACHIDHG 180
 QY 181 KTVKGTQKLRVGLFKPKFQDGGRGANDSTAMPEQMKFQIRVKTHTVPKGIKGNAND 240
 Db 181 KTVKGTQKLRVGLFKPKFQDGGRGANDSTAMPEQMKFQIRVKTHTVPKGIKGNAND 240
 QY 241 IGMDYVALLLELKKPKRKNKIGVSPPAKQLPGGRIFHSGYNDNDRPGLNLYVYRFDVKDE 300
 Db 241 IGMDYVALLLELKKPKRKNKIGVSPPAKQLPGGRIFHSGYNDNDRPGLNLYVYRFDVKDE 300
 QY 301 TYDLLVOCCDAOPGASGVYVVMWKQKQKWKRIIGIESGHQWVDMNGSPQDENVAVR 360
 Db 301 TYDLLVOCCDAOPGASGVYVVMWKQKQKWKRIIGIESGHQWVDMNGSPQDENVAVR 360
 QY 361 ITPKVAQICYWIKNYLDREG 383
 Db 361 ITPKVAQICYWIKNYLDREG 383
 RESULT 12.
 AAU29048
 ID AAU29048 standard; protein; 383 AA.
 XX
 AC AAU29048;
 XX
 DT 18-DEC-2001 (first entry)
 XX
 DE Human PRO polypeptide sequence #25.
 XX
 KW PRO polypeptide; mammal; tumour; cancer; human; cattle; horse; sheep;

KW dog; cat; pig; goat; rabbit; tumour necrosis factor alpha; TNF-alpha;
 KW blood; chondrocyte cell; cell proliferation; cell differentiation; colon;
 KW adrenal; lung; breast; prostate; rectum; cervix; liver; genetic disorder.
 XX
 OS Homo sapiens.
 XX
 XX WO200168848-A2.
 XX
 XX 20-SEP-2001.
 XX
 XX 28-FEB-2001; 2001WO-US006520.
 XX
 XX 01-MAR-2000; 2000WO-US005601.
 XX 02-MAR-2000; 2000WO-US005841.
 XX 03-MAR-2000; 2000US-0187202P.
 XX 06-MAR-2000; 2000US-0186968P.
 XX 14-MAR-2000; 2000US-0189320P.
 XX 14-MAR-2000; 2000US-0189328P.
 XX 15-MAR-2000; 2000WO-US006884.
 XX 21-MAR-2000; 2000US-0190828P.
 XX 21-MAR-2000; 2000US-0191007P.
 XX 21-MAR-2000; 2000US-0191048P.
 XX 21-MAR-2000; 2000US-0191314P.
 XX 28-MAR-2000; 2000US-0192655P.
 XX 29-MAR-2000; 2000US-0193032P.
 XX 29-MAR-2000; 2000US-0193053P.
 XX 30-MAR-2000; 2000WO-US008439.
 XX 04-APR-2000; 2000US-0194449P.
 XX 04-APR-2000; 2000US-0194647P.
 XX 11-APR-2000; 2000US-0195975P.
 XX 11-APR-2000; 2000US-0196000P.
 XX 11-APR-2000; 2000US-0196187P.
 XX 11-APR-2000; 2000US-0196690P.
 XX 11-APR-2000; 2000US-0196820P.
 XX 18-APR-2000; 2000US-0198121P.
 XX 18-APR-2000; 2000US-0198585P.
 XX 25-APR-2000; 2000US-0199397P.
 XX 25-APR-2000; 2000US-0199550P.
 XX 25-APR-2000; 2000US-0199654P.
 XX 03-MAY-2000; 2000US-0201516P.
 XX 17-MAY-2000; 2000WO-US013705.
 XX 22-MAY-2000; 2000WO-US014042.
 XX 30-MAY-2000; 2000WO-US014941.
 XX 02-JUN-2000; 2000WO-US015264.
 XX 05-JUN-2000; 2000US-0209832P.
 XX 28-JUL-2000; 2000WO-US020710.
 XX 22-AUG-2000; 2000US-00644848.
 XX 24-AUG-2000; 2000WO-US023328.
 XX 08-NOV-2000; 2000WO-US030952.
 XX 01-DEC-2000; 2000WO-US032678.
 XX 20-DEC-2000; 2000WO-US034956.
 XX
 (GETH) GENENTECH INC.
 PA
 XX Baker KP, Chen J, Desnoyers L, Goddard A, Godowski PJ, Gurney AL;
 PI Pan J, Smith V, Watanabe CK, Wood WI, Zhang Z;
 XX
 XX WPI; 2001-602746/68.
 XX N-PSDB; AAS45949.
 XX
 XX Novel nucleic acids encoding PRO polypeptides, used to diagnose the
 PT presence of tumors, such as prostate and breast tumors, in mammals and to
 PT screen for modulators of the compounds.
 XX
 XX Claim 11; Fig 50; 774pp; English.
 PS
 XX Sequences AAU29024-AAU29328 represent PRO polypeptides of the invention.
 CC The PRO polypeptides and their associated nucleic acids can be used to
 CC detect the presence of a tumour in a mammal by comparing the level of
 CC expression of a PRO polypeptide in a test sample of cells from the animal
 CC and a control sample of normal cells, whereby a higher level of
 CC expression in the test sample indicates the presence of a tumour in the
 CC mammal. Mammals include dogs, cats, cattle, horses, sheep, pigs, goats

CC and rabbits but are preferably human. The polypeptides can be used to
CC stimulate tumour necrosis factor (TNF) alpha release from human blood,
CC when contacted with it. A specific polypeptide can be used to stimulate
CC the proliferation or differentiation of chondrocyte cells. The PRO
CC proteins can be used to determine the presence of tumours and also
CC susceptibility to tumour development, particularly adrenal, lung, colon,
CC breast, prostate, rectal, cervical, or liver tumours, in mammalian
CC subjects. The oligonucleotide probes specific for the PRO nucleic acids
CC can be used for genetic analysis of individuals with genetic disorders
XX

SQ Sequence 383 AA;

Query Match	100.0%;	Score 2080;	DB 4;	Length 383;
Best Local Similarity	100.0%;	Pred. NO. 1.8e-145;		
Matches 383;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;

Qy	1	MAGTGLLFLFLLCVAGQVSPSPAPKPTWPAYRLPVVLPOSTLNLAKEPFGAEAKLE	60
Db	1	MAGTGLLFLFLLCVAGQVSPSPAPKPTWPAYRLPVVLPOSTLNLAKEPFGAEAKLE	60
Qy	61	VSSSCGPGCHKGTPLPYBEAKQYLSYETILYANGSRTEQTQVGIYILSSSGDGAQHRDGS	120
Db	61	VSSSCGPGCHKGTPLPYBEAKQYLSYETILYANGSRTEQTQVGIYILSSSGDGAQHRDGS	120
Qy	121	SGKSRKRQIYGVDSRPSIFGKDFLLNYPSTSVKLSCTGCTGLVAEKHVLTAACHIDG	180
Db	121	SGKSRKRQIYGVDSRPSIFGKDFLLNYPSTSVKLSCTGCTGLVAEKHVLTAACHIDG	180
Qy	181	KTVKGTQKLRVGLFKPKFGKGGRGANDSTAMPEQMKFQWIRKRVKTHVPKGIKGNAND	240
Db	181	KTVKGTQKLRVGLFKPKFGKGGRGANDSTAMPEQMKFQWIRKRVKTHVPKGIKGNAND	240
Qy	241	IGMDYDVALLELKKPKRKFVKIGVSPPAKQLPGRIHFSGYDNDPFGNLVYRFDVKDE	300
Db	241	IGMDYDVALLELKKPKRKFVKIGVSPPAKQLPGRIHFSGYDNDPFGNLVYRFDVKDE	300
Qy	301	TYDLLYQCCDAQPGASGSGVYVRMWKQKQKWERKIIGIFSGHGWDMNGSPQDFNVAVR	360
Db	301	TYDLLYQCCDAQPGASGSGVYVRMWKQKQKWERKIIGIFSGHGWDMNGSPQDFNVAVR	360
Qy	361	ITPLKVAQICWIKGNVLDREG 383	
Db	361	ITPLKVAQICWIKGNVLDREG 383	

RESULT 13

ABU58424

ID ABU58424 standard; protein; 383 AA.

XX AC ABU58424;

XX XX 15-APR-2003 (first entry)

XX XX Human PRO polypeptide #25.

XX Human; PRO; cytostatic; tumour; cancer; breast; lung; stomach; liver;
KW dog; cat; cow; horse; sheep; pig; goat; rabbit; ADEPT;
KW antibody-dependent enzyme mediated prodrug therapy.

OS Homo sapiens.

XX XX US2003027272-A1.

XX XX 06-FEB-2003.

XX XX 21-JUN-2002; 2002US-00176492.

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XX OS
XX Homo sapiens.
XX US2003032127-A1.
XX 13-FEB-2003.
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PR	02-OCT-1998;	98US-0102955P.
PR	06-OCT-1998;	98US-0103258P.

Query Match	100.0%;	Score 2080;	DB 6;	Length 383;
Best Local Similarity	100.0%;	Pred. No. 1.8e-145;		
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Qy	1	MAGIPGLLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPPOSTLNLAKPDPFGAAKLE	60	
Db	1	MAGIPGLLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPPOSTLNLAKPDPFGAAKLE	60	
Qy	61	VSSSCGPOCHKGTPPTYEEAKQYLSYETLYANGSRTETQVGIYIILSSSGDGAQHRDGS	120	
Db	61	VSSSCGPOCHKGTPPTYEEAKQYLSYETLYANGSRTETQVGIYIILSSSGDGAQHRDGS	120	
Qy	121	SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEKHVLTAACHIDG	180	
Db	121	SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEKHVLTAACHIDG	180	
Qy	181	KTYVKGTOKLRYGFLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGWIKGNAND	240	
Db	181	KTYVKGTOKLRYGFLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGWIKGNAND	240	
Qy	241	IGMDYDYLLELKKPHKRFKMGKIGVSPPAKQLPGGRIFHFGYDNDNRPGNLVYRFCVDKDE	300	
Db	241	IGMDYDYLLELKKPHKRFKMGKIGVSPPAKQLPGGRIFHFGYDNDNRPGNLVYRFCVDKDE	300	
Qy	301	TYDLLYQOCDAOPGASGSGVYVMMKROQOKWERKIIGIFSGHQWYDMNGSPQDFNVAVR	360	
Db	301	TYDLLYQOCDAOPGASGSGVYVMMKROQOKWERKIIGIFSGHQWYDMNGSPQDFNVAVR	360	
Qy	361	ITPLKYAQICYWIKGNLYDCREG	383	
Db	361	ITPLKYAQICYWIKGNLYDCREG	383	

Search completed: July 1, 2005, 21:03:29
Job time : 99.1294 secs

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: July 1, 2005, 20:57:14 ; Search time 25.9272 Seconds
(without alignments)
1102.727 Million cell updates/sec

Title: US-09-658-677-18
Perfect score: 2080
Sequence: 1 MAGIPGLFLFLFLLCAVGQ.....LKVAQICYWIKGNVLDREG 383

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 513545 seqs, 74649064 residues

Total number of hits satisfying chosen parameters: 513545

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued Patents AA:*
1: /cgn2_6/ptodata/1/iaa/5A COMB.pep.*
2: /cgn2_6/ptodata/1/iaa/5B COMB.pep.*
3: /cgn2_6/ptodata/1/iaa/6A COMB.pep.*
4: /cgn2_6/ptodata/1/iaa/6B COMB.pep.*
5: /cgn2_6/ptodata/1/iaa/PCTUS COMB.pep.*
6: /cgn2_6/ptodata/1/iaa/backfiles1.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	2080	100.0	383	3	US-09-072-384-18
2	2080	100.0	383	4	US-09-907-794A-261
3	2080	100.0	383	4	US-09-905-125A-261
4	2080	100.0	383	4	US-09-902-775A-261
5	2080	100.0	383	4	US-09-906-700-261
6	2080	100.0	383	4	US-09-903-603A-261
7	2080	100.0	383	4	US-09-904-920A-261
8	2080	100.0	383	4	US-09-909-064-261
9	2080	100.0	383	4	US-09-905-381A-261
10	2080	100.0	383	4	US-09-906-618-261
11	2080	100.0	392	3	US-09-072-384-15
12	2044	98.3	392	3	US-09-072-384-2
13	202.5	9.7	314	4	US-09-551-826D-6
14	193	9.3	316	4	US-09-551-826D-2
15	177	8.5	222	1	US-08-090-048-1
16	177	8.5	222	2	US-08-292-550-1
17	177	8.5	222	2	US-07-927-661A-1
18	176.5	8.5	318	4	US-09-551-826D-10
19	140.5	6.8	239	4	US-09-107-433-3471
20	135.5	6.5	313	4	US-09-551-826D-14
21	132	6.3	218	4	US-09-583-110-3950
22	129.5	6.2	302	4	US-09-551-826D-12
23	129	6.2	433	4	US-09-949-016-8220
24	127	6.1	303	4	US-09-551-826D-8
25	127	6.1	320	4	US-09-489-039A-13989
26	126.5	6.1	356	4	US-09-902-540-12881
27	122.5	5.9	238	4	US-09-664-595A-15

Sequence 89, Appl
Sequence 89, Appl
Sequence 89, Appl
Sequence 89, Appl
Sequence 89, Appl
Sequence 32, Appl
Sequence 89, Appl
Sequence 89, Appl
Sequence 7, Appl
Sequence 7, Appl
Sequence 7, Appl
Sequence 2, Appl
Sequence 59, Appl
Sequence 44, Appl
Patent No. 5223425
Patent No. 5223425
Patent No. 5223425

28 122 5.9 256 3 US-08-306-769-89
29 122 5.9 256 3 US-08-306-616-89
30 122 5.9 256 3 US-08-817-795-89
31 122 5.9 256 3 US-08-639-075A-89
32 122 5.9 256 3 US-09-012-431-89
33 122 5.9 256 3 US-09-032-215-32
34 122 5.9 256 3 US-09-012-692-89
35 122 5.9 256 3 US-08-306-613-89
36 122 5.9 256 5 PCT-US95-14442A-89
37 120 5.8 284 4 US-09-387-375-7
38 120 5.8 284 4 US-10-041-400A-7
39 120 5.8 284 4 US-10-042-091A-7
40 117 5.6 437 1 US-08-487-037-2
41 115.5 5.6 241 3 US-08-944-483-59
42 114 5.5 228 3 US-08-944-483-44
43 114 5.5 253 6 5223425-8
44 114 5.5 253 6 5223425-8
45 113 5.4 238 6 5223425-5

ALIGNMENTS

RESULT 1
US-09-072-384-18
; Sequence 18, Application US/09072384
; Patent No. 6153420
; GENERAL INFORMATION:
; APPLICANT: Sheppard, Paul O.
; TITLE OF INVENTION: SERINE PROTEASE POLYPEPTIDES
; AND MATERIALS AND METHODS FOR MAKING THEM
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSSEE: ZymoGenetics, Inc.
; STREET: 1201 Eastlake Avenue East
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98102
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/072,384
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Parker, Gary E
; REGISTRATION NUMBER: 31,648
; REFERENCE/DOCKET NUMBER: 97-16C1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-442-6673
; TELEFAX: 206-442-6678
; TELEX:
; INFORMATION FOR SEQ ID NO: 18:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 383 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FRAGMENT TYPE: internal
; FEATURE:
; NAME/KEY: Signal Sequence
; LOCATION: 1...19
; OTHER INFORMATION:
; US-09-072-384-18

Query Match	100.0%	Score 2080;	DB 3;	Length 383;
Best Local Similarity	100.0%;	Pred. No. 5.4e-220;		
Matches 383;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
Qy	1	MAGIPGGLFLLFFLLCAVGQVSPYSAPWKPTWPAYRLPVVLPOSTLNLAKPDFGAEAKLE	60	
Db	1	MAGIPGGLFLLFFLLCAVGQVSPYSAPWKPTWPAYRLPVVLPOSTLNLAKPDFGAEAKLE	60	
Qy	61	VSSCGPQCHKGTPLPTYEEAKQYLSYETLYANGSRSTETQVGIYILSSSGDGAQHRDGS	120	
Db	61	VSSCGPQCHKGTPLPTYEEAKQYLSYETLYANGSRSTETQVGIYILSSSGDGAQHRDGS	120	
Qy	121	SGSRRRKQIYGYDSRFSIIFGKDFLLNYPFSTSVKLSGTGCTTLVAEKHVLTAACHIDG	180	
Db	121	SGSRRRKQIYGYDSRFSIIFGKDFLLNYPFSTSVKLSGTGCTTLVAEKHVLTAACHIDG	180	
Qy	181	KTVYKGTQKLRVGLPKPFKFDGGRGANDSAMSPEQMKFQWRVKRTHVPKGIWKGNAND	240	
Db	181	KTVYKGTQKLRVGLPKPFKFDGGRGANDSAMSPEQMKFQWRVKRTHVPKGIWKGNAND	240	
Qy	241	IGMDYDVALLELKKPHRKPKWKIGVSPPAKQLPGRIHFSGYDNDNRGNLVYRCDVKDE	300	
Db	241	IGMDYDVALLELKKPHRKPKWKIGVSPPAKQLPGRIHFSGYDNDNRGNLVYRCDVKDE	300	
Qy	301	TYDLLYQOCDAQPGASGSGVYVRMWKQOQKWERKIIGIFSGHOWDMNGSPQDFNVAVR	360	
Db	301	TYDLLYQOCDAQPGASGSGVYVRMWKQOQKWERKIIGIFSGHOWDMNGSPQDFNVAVR	360	
Qy	361	ITPLKYAQICYWKGNLYDCREG	383	
Db	361	ITPLKYAQICYWKGNLYDCREG	383	

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RESULT 2
US-09-907-794A-261
; Sequence 261, Application US/09907794A
; Patent No. 6635468
;
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, AVI
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
;
; TITLE OF INVENTION: Secreted and Transmembrane Proteins
; FILE REFERENCE: Acids Encoding th
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907794A
; CURRENT FILING DATE: 2001-07-17
; PRIOR APPLICATION NUMBER: PCT/US00/044
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143, 0
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145, 6

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? PRIOR FILING DATE: 1999-07-26
? PRIOR APPLICATION NUMBER: US 60/146,222
? PRIOR FILING DATE: 1999-07-28
? PRIOR APPLICATION NUMBER: PCT/US99/20594
? PRIOR FILING DATE: 1999-09-08
? PRIOR APPLICATION NUMBER: PCT/US99/20944
? PRIOR FILING DATE: 1999-09-13
? PRIOR APPLICATION NUMBER: PCT/US99/21090
? PRIOR FILING DATE: 1999-09-15
? PRIOR APPLICATION NUMBER: PCT/US99/21547
? PRIOR FILING DATE: 1999-09-15
? PRIOR APPLICATION NUMBER: PCT/US99/23089
? PRIOR FILING DATE: 1999-10-05
? PRIOR APPLICATION NUMBER: PCT/US99/28214
? PRIOR FILING DATE: 1999-11-29
? PRIOR APPLICATION NUMBER: PCT/US99/28313
? PRIOR FILING DATE: 1999-11-30
? PRIOR APPLICATION NUMBER: PCT/US99/28564
? PRIOR FILING DATE: 1999-12-02
? PRIOR APPLICATION NUMBER: PCT/US99/28565
? PRIOR FILING DATE: 1999-12-02
? PRIOR APPLICATION NUMBER: PCT/US99/30095
? PRIOR FILING DATE: 1999-12-16
? PRIOR APPLICATION NUMBER: PCT/US99/30911
? PRIOR FILING DATE: 1999-12-20
? PRIOR APPLICATION NUMBER: PCT/US99/30999
? PRIOR FILING DATE: 1999-12-20
? PRIOR APPLICATION NUMBER: PCT/US00/00219
? PRIOR FILING DATE: 2000-01-05
? NUMBER OF SEQ ID NOS: 423
? SEQ ID NO 261
? LENGTH: 383
? TYPE: PRT
? ORGANISM: Homo Sapien
? US-09-907-794A-261

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Query Match	100.0%;	Score 2080;	DB 4;	Length 383;
Best Local Similarity	100.0%;	Pred. No. 5.4e-220;		
Matches 383;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
Qy	1	MAGIPGLLFLFLLC	AVGVSPYSAPWKPTW	PAYRLPVVLPOSTLNLAKDFGAEAKLE 60
Db	1	MAGIPGLLFLFLLC	AVGVSPYSAPWKPTW	PAYRLPVVLPOSTLNLAKDFGAEAKLE 60
Qy	61	VSSCGPQCHKGTPL	PYBEAKQYLSYETIY	ANGSRSTETQVGIYILSSSGDGAOHRDGS 120
Db	61	VSSCGPQCHKGTPL	PYBEAKQYLSYETIY	ANGSRSTETQVGIYILSSSGDGAOHRDGS 120
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Db	121	SGKRRRKQIYGVDS	RSIFGKQFLNYPST	SVKLTGCTGTVAEKHVLTAAHCIDHG 180
Qy	181	KTVYKGTQKLVRG	FLKPKFDGGRGAND	STSAMPEQMKFQIRVKRTHVPKGIWKGAND 240
Db	181	KTVYKGTQKLVRG	FLKPKFDGGRGAND	STSAMPEQMKFQIRVKRTHVPKGIWKGAND 240
Qy	241	IGMDYDALLLEL	KPKHKPKMGIVS	PPAKQLPGRIHFSGYDNDRCNLVYRCDVKDE 300
Db	241	IGMDYDALLLEL	KPKHKPKMGIVS	PPAKQLPGRIHFSGYDNDRCNLVYRCDVKDE 300
Qy	301	TYDLLYQQCDAQ	PCGASGSGVYVRMW	KRQQQKWERKIIGIFSGHQMVMNMGSPQDFNVAVR 360
Db	301	TYDLLYQQCDAQ	PCGASGSGVYVRMW	KRQQQKWERKIIGIFSGHQMVMNMGSPQDFNVAVR 360
Qy	361	ITPLKYAQICYWI	KGNLYDCREG	383
Db	361	ITPLKYAQICYWI	KGNLYDCREG	383

RESULT 3
US-09-905-125A-261
; Sequence 261, Application US/09905125A
; Patent No. 6664376

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; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gernitsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,125A
; PRIOR FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/145,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-905-125A-261
; Query Match 100.0%; Score 2080; DB 4; Length 383;

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Best Local Similarity 100.0%; Pred. No. 5.4e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGIPGLLLFLFLCAVGVSPYSAPWKTWPAYRLPVVLPOSTLNLAKPDPFGAEAKLE 60
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 1 MAGIPGLLLFLFLCAVGVSPYSAPWKTWPAYRLPVVLPOSTLNLAKPDPFGAEAKLE 60
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |

Qy 61 VSSCGPQCHKGTPPTYEAKOYLSYETLYANGSRRTETQVGIYILSSSGDGAQHRDGS 120
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 61 VSSCGPQCHKGTPPTYEAKOYLSYETLYANGSRRTETQVGIYILSSSGDGAQHRDGS 120
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |

Qy 121 SGKRRKRIQYGYDSRFSIFGKDFLLNYPSTSVKLTGTCTGLVAEKHVLTAACHIDG 180
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 121 SGKRRKRIQYGYDSRFSIFGKDFLLNYPSTSVKLTGTCTGLVAEKHVLTAACHIDG 180
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |

Qy 181 KTVYKGTQKLRVGFLLPKPKDGGRGANDSTSAPEQMKFQWIRVKTHTVPKGTWKNAND 240
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 181 KTVYKGTQKLRVGFLLPKPKDGGRGANDSTSAPEQMKFQWIRVKTHTVPKGTWKNAND 240
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |

Qy 241 IGMDDYVALLLEKPKHKKFMKIGVSPKQKPGRIHFGSYDNDPGLNLYVRFCDVKDE 300
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 241 IGMDDYVALLLEKPKHKKFMKIGVSPKQKPGRIHFGSYDNDPGLNLYVRFCDVKDE 300
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |

Qy 301 TYDLLYQCCDAQPCASGSGVYVVMWKRQQQKWKRIIGFSGHQMVDNMGSPQDFNVAVR 360
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 301 TYDLLYQCCDAQPCASGSGVYVVMWKRQQQKWKRIIGFSGHQMVDNMGSPQDFNVAVR 360
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |

Qy 361 ITPKVAQICYWIKNYLDCREG 383
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 361 ITPKVAQICYWIKNYLDCREG 383
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |

RESULT 4
US-09-902-775A-261
; Sequence 261, Application US/09902775A
; Patent No. 6686451
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gernitsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/902,775A
; CURRENT FILING DATE: 2001-07-10
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26

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; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
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; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-906-700-261

Query Match 100.0%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 5.4e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGTIGLLFLFLLCAVGVSPYAPWKTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
Db 1 MAGTIGLLFLFLLCAVGVSPYAPWKTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60

Qy 61 VSSSCGQCHKGTPLPYBAKQVLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS 120
Db 61 VSSSCGQCHKGTPLPYBAKQVLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS 120

Qy 121 SGKSRKRQIYGVDSRESIFGKDFLLNYPSTSVKLSCTGCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGVDSRESIFGKDFLLNYPSTSVKLSCTGCTGLVAEKHVLTAACHIDG 180

Qy 181 KTVYVKGTKLRVGLPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240
Db 181 KTVYVKGTKLRVGLPKFKDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240

Qy 241 IGMDDYDVALLELKKPHRKKFMKIGVSPPAKQLPGRIFHSGYNDNRPGLNLYRFDVKDE 300
Db 241 IGMDDYDVALLELKKPHRKKFMKIGVSPPAKQLPGRIFHSGYNDNRPGLNLYRFDVKDE 300

Qy 301 TYDLLYQCCDAQPGASGVYVRMKQOQKWERKIIGFSGHQWDMGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAQPGASGVYVRMKQOQKWERKIIGFSGHQWDMGSPQDFNVAVR 360

Qy 361 ITPLKYAQICWIKGNLYDCREG 383
Db 361 ITPLKYAQICWIKGNLYDCREG 383

RESULT 5
US-09-906-700-261
; Sequence 261, Application US/09906700
; Patent No. 6723535
; GENERAL INFORMATION:

; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/906,700
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-906-700-261

Query Match 100.0%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 5.4e-220;

Matches	383;	Conservative	0;	Mismatches	0;	Indels	0;	Gaps	0;
1	MAGIPGLLFLFLFFLLCAVQGVSPYSAPWKPTWPAYRLPVLPQSTLNIAKPDFGAEAKLE	60							
1	MAGIPGLLFLFLFFLLCAVQGVSPYSAPWKPTWPAYRLPVLPQSTLNIAKPDFGAEAKLE	60							
61	VSSCGPQCHKGTPLPITYEEAAQKYLSEYTLVANGSRSTETQVGIYILSSSGDGAQHRDGS	120							
61	VSSCGPQCHKGTPLPITYEEAAQKYLSEYTLVANGSRSTETQVGIYILSSSGDGAQHRDGS	120							
121	SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTGTLVAEKHVLTAACHIDG	180							
121	SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTGTLVAEKHVLTAACHIDG	180							
181	KTYVKGTKQLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND	240							
181	KTYVKGTKQLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND	240							
241	IGMDYDYALLBKPKHKRKFWKIIGVSPPAKLPQGRIFHSGYDNDRPGNLVYRFCDVKDE	300							
241	IGMDYDYALLBKPKHKRKFWKIIGVSPPAKLPQGRIFHSGYDNDRPGNLVYRFCDVKDE	300							
301	TYDLLYQQCDAPGASGSGVYVRWVKRQQQKWERKIIIPFSGHQVDMNGSPQDFNVAVR	360							
301	TYDLLYQQCDAPGASGSGVYVRWVKRQQQKWERKIIIPFSGHQVDMNGSPQDFNVAVR	360							
361	ITPLKYAQICYWIKNYLDCREG	383							
361	ITPLKYAQICYWIKNYLDCREG	383							

RESULT 6
US-09-903-603A-261

: Sequence 261. Application US/09903603A

: Patent No. 6767995
; Sequence 201, Application 05/0330303A

; PATENT NO. 6767995
; GENERAL INFORMATION.

GENERAL INFORMATION:

APPLICANT: Genentech, Inc.

; APPLICANT: Ashkenazi, Avi

APPLICANT: Botstein, David

APPLICANT: Desnoyers, Luc

APPLICANT: Desmoyers, Luc
APPLICANT: Eaton, Dan L.

APPLICANT: EATON, DAN L.
APPLICANT: Ferrara, Napolitano

APPLICANT: Ferrara, Napoleone
APPLICANT: Ellsworth, Ellen

APPLICANT: Filvaroff, Ellen

APPLICANT: Fong, Sherman

; APPLICANT: Gao, Wei-Qiang

APPLICANT: Gerber, Hanspeter

APPLICANT: Gerritsen, Mary E.

APPLICANT: Gottschen, Mary E.
APPLICANT: Goddard, A.

APPLICANT: GODDARD, A.
APPLICANT: Godowski Paul J.

APPLICANT: GODOWSKI, PAUL J.
APPLICANT: CATALDI, CHRISTOPHER J.

APPLICANT: Grimaldi, Christopher J.

APPLICANT: Gurney, Austin L.

; APPLICANT: Hillan, Kenneth, J.

APPLICANT: Kljavin, Ivar J.

APPLICANT: Mather, Jennie P.

APPLICANT: pan. James

APPLICANT: PAN, JAMES
APPLICANT: PAN, NICHOLAS F

APPLICANT: PAONI, Nicholas F.
APPLICANT: PAONI, Nicholas F.

APPLICANT: Roy, Margaret Ann

APPLICANT: Stewart, Timothy A.

APPLICANT: Tumas, Daniel

APPLICANT: Williams, P. Mickey

APPLICANT: Wood, William. I.

1. TITLE OF INVENTION: Secreted and Trans

; TITLE OF INVENTION: secreted and transcribed RNA

; TITLE OF INVENTION: Acids Encoding the

; FILE REFERENCE: GNE.1618P2C12

; CURRENT APPLICATION NUMBER: US/09/903,

; CURRENT FILING DATE: 2001-07-11

PRIOR APPLICATION NUMBER: PCT/US00/044

PRIOR FILING DATE: 2000-02-22

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; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-903-603A-261

Query Match      100.0%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 5.4e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy  1  MAGIPGLLFLFLLLCAVGVQSPYSAPWKPTWPAYRLPVVLPSTLNLAKPDPFGAEAKLE 60
Db  1  MAGIPGLLFLFLLLCAVGVQSPYSAPWKPTWPAYRLPVVLPSTLNLAKPDPFGAEAKLE 60

Qy  61  VSSSCGPOCHKGTPLPTYBEAKQYLSYETLYANGSRRTETQVGIIYILSSSGDGAQHRDGS 120
Db  61  VSSSCGPOCHKGTPLPTYBEAKQYLSYETLYANGSRRTETQVGIIYILSSSGDGAQHRDGS 120

Qy  121  SGKSRKRQIYGVDSRFSIFGKDFLNYPFSTSVKLSCTGCTGTILVAEKHVLTAACHIDG 180
Db  121  SGKSRKRQIYGVDSRFSIFGKDFLNYPFSTSVKLSCTGCTGTILVAEKHVLTAACHIDG 180

Qy  181  KTVVKGTTQKLRVGFLKPKFGDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGMWIKGNAND 240
Db  181  KTVVKGTTQKLRVGFLKPKFGDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGMWIKGNAND 240

Qy  241  IGMDDYDYLLELKKPKHKRFKMIKGVSPPAKQLPGGRIHFSGYNDREPNLVYRFDVKDE 300
Db  241  IGMDDYDYLLELKKPKHKRFKMIKGVSPPAKQLPGGRIHFSGYNDREPNLVYRFDVKDE 300

Qy  301  TYDLLYQQCDAQPGAGSGSGVYVMWKRQOQKWERKIIGIPSGHQWVDMNGSPQDFNVAVR 360
Db  301  TYDLLYQQCDAQPGAGSGSGVYVMWKRQOQKWERKIIGIPSGHQWVDMNGSPQDFNVAVR 360

Qy  361  ITPKVAQICYMIKGNLYDCREG 383
Db  361  ITPKVAQICYMIKGNLYDCREG 383

RESULT 7
US-09-904-920A-261
; Sequence 261, Application US/09904920A
; Patent No. 6806352
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.

```

APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/904,920A
PRIOR FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28364
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/30095
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-904-920A-261
Query Match 100.0%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 5.4e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLELLFLLCAVGVQVSPYSAPWKPTWPAYRLPVVLPOSTLNLAKPFGAEAKLE 60
DB 1 MAGIPGLLELLFLLCAVGVQVSPYSAPWKPTWPAYRLPVVLPOSTLNLAKPFGAEAKLE 60
QY 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTETQVGIYIILSSGSGAQHRDGS 120
DB 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTETQVGIYIILSSGSGAQHRDGS 120
QY 121 SGKSRKROIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEKHVLTAACHIDG 180
DB 121 SGKSRKROIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEKHVLTAACHIDG 180
QY 181 KTVVKGTKLRVGFLLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
DB 181 KTVVKGTKLRVGFLLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
QY 241 IGMDYDYLLELKKPHKRFKMGIVSPPAKQLPGGRIHFGSYDNDPRGNLVYRFDVKDE 300
DB 241 IGMDYDYLLELKKPHKRFKMGIVSPPAKQLPGGRIHFGSYDNDPRGNLVYRFDVKDE 300
QY 301 TYDLLVQOCDAOPGASGSGVYVVMWKRQCKWKRIIGIFSGHGWDMNGSPQDFNVAVR 360
DB 301 TYDLLVQOCDAOPGASGSGVYVVMWKRQCKWKRIIGIFSGHGWDMNGSPQDFNVAVR 360
QY 361 ITPLKYAQICYMIKGNLYDCREG 383
DB 361 ITPLKYAQICYMIKGNLYDCREG 383
RESULT 8
US-09-909-064-261
Sequence 261, Application US/09909064
Patent No. 6818449
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/909,064
PRIOR FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28

APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/905,381A
CURRENT FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/30095
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-909-064-261

Query Match 100.0%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 5.4e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAGGVSPVSPAPKPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLFLFLLCAGGVSPVSPAPKPTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPGCHGKPTLPYBEAKQVLSYETLYANGSRRTETQVGIYILSSGSGAQHRDGS 120
DB 61 VSSSCGPGCHGKPTLPYBEAKQVLSYETLYANGSRRTETQVGIYILSSGSGAQHRDGS 120

QY 121 SGSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLTGCTGTVAEKHVLTAACHIDG 180
DB 121 SGSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLTGCTGTVAEKHVLTAACHIDG 180

QY 181 KTVVGTQKLVGFLKPKFVGGGANDSTSAMPEQMKFQWIRKTRHVPKGIKGNAND 240
DB 181 KTVVGTQKLVGFLKPKFVGGGANDSTSAMPEQMKFQWIRKTRHVPKGIKGNAND 240

QY 241 IGMDDYVALLLEKKPKRKFEMKIGVSPAPKQALPGRIHFSGYDNDRCNLVYRFDVKDE 300
DB 241 IGMDDYVALLLEKKPKRKFEMKIGVSPAPKQALPGRIHFSGYDNDRCNLVYRFDVKDE 300

QY 301 TYDLLYQQCDAQPGASGSGVYVVRWKRFQOQKWERKIIGIFSGHQQWDMGSPQDFNVAVR 360
DB 301 TYDLLYQQCDAQPGASGSGVYVVRWKRFQOQKWERKIIGIFSGHQQWDMGSPQDFNVAVR 360

QY 361 ITPKYAQICYIKGNVLDREG 383
DB 361 ITPKYAQICYIKGNVLDREG 383

RESULT 9
US-09-905-381A-261
Sequence 261, Application US/09905381A
Patent No. 6818746
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi

Query Match 100.0%; Score 2080; DB 4; Length 383;
Best Local Similarity 100.0%; Pred. No. 5.4e-220;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
Qy 61 VSSSCGQCHKGTPLTPEAKOVLSTYANGSRTEQVGIYILSSSGDGAQHRDGS 120
Db 61 VSSSCGQCHKGTPLTPEAKOVLSTYANGSRTEQVGIYILSSSGDGAQHRDGS 120
Qy 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEKHVLTAACHIDG 180
Db 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEKHVLTAACHIDG 180
Qy 181 KTVYVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
Db 181 KTVYVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
Qy 241 IGMDYDVALLELKKPKHKKFKWKICVSPAKOLPGRIHFGSGYDNDPRGNLVYRFDVKDE 300
Db 241 IGMDYDVALLELKKPKHKKFKWKICVSPAKOLPGRIHFGSGYDNDPRGNLVYRFDVKDE 300
Qy 301 TYDLYQQCDAQPGASGSGVYVVMWKRQQQKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Db 301 TYDLYQQCDAQPGASGSGVYVVMWKRQQQKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Qy 361 ITPLKVAQICYWIKNYLDCREG 383
Db 361 ITPLKVAQICYWIKNYLDCREG 383

RESULT 10

US-09-906-618-261

; Sequence 261, Application US/09906618

; Patent No. 6828146

; GENERAL INFORMATION:

; APPLICANT: Genentech, Inc.

; APPLICANT: Ashkenazi, Avi

; APPLICANT: Botstein, David

; APPLICANT: Desnoyers, Luc

; APPLICANT: Eaton, Dan L.

; APPLICANT: Ferrara, Napoleone

; APPLICANT: Filvaroff, Ellen

; APPLICANT: Fong, Sherman

; APPLICANT: Gao, Wei-Qiang

; APPLICANT: Gerber, Hanspeter

; APPLICANT: Gerritsen, Mary E.

; APPLICANT: Goddard, A.

; APPLICANT: Godowski, Paul J.

; APPLICANT: Grimaldi, Christopher J.

; APPLICANT: Gurney, Austin L.

; APPLICANT: Hillan, Kenneth, J.

; APPLICANT: Kijavlin, Ivar J.

; APPLICANT: Mather, Jennie P.

; APPLICANT: Pan, James

; APPLICANT: Paoni, Nicholas F.

; APPLICANT: Roy, Margaret Ann

; APPLICANT: Stewart, Timothy A.

; APPLICANT: Tumas, Daniel

; APPLICANT: Williams, P. Mickey

; APPLICANT: Wood, William, I.

; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic

; FILE REFERENCE: 10466-14

; CURRENT APPLICATION NUMBER: US/09/906,618

; CURRENT FILING DATE: 2001-07-16

; PRIOR APPLICATION NUMBER: PCT/US00/04414

; PRIOR FILING DATE: 2000-02-22

; PRIOR APPLICATION NUMBER: US 60/143,048

; PRIOR FILING DATE: 1999-07-07

; PRIOR APPLICATION NUMBER: US 60/145,698

; PRIOR FILING DATE: 1999-07-26

; PRIOR APPLICATION NUMBER: US 60/146,222

; PRIOR FILING DATE: 1999-07-28

; PRIOR APPLICATION NUMBER: PCT/US99/20594

; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-906-618-261

Query Match 100.0%; Score 2080; DB 4; Length 383;

Best Local Similarity 100.0%; Pred. No. 5.4e-220;

Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPSTLNLAKEPFGAEAKLE 60
Qy 61 VSSSCGQCHKGTPLTPEAKOVLSTYANGSRTEQVGIYILSSSGDGAQHRDGS 120
Db 61 VSSSCGQCHKGTPLTPEAKOVLSTYANGSRTEQVGIYILSSSGDGAQHRDGS 120
Qy 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEKHVLTAACHIDG 180
Db 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLSCTGTLVAEKHVLTAACHIDG 180
Qy 181 KTVYVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
Db 181 KTVYVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
Qy 241 IGMDYDVALLELKKPKHKKFKWKICVSPAKOLPGRIHFGSGYDNDPRGNLVYRFDVKDE 300
Db 241 IGMDYDVALLELKKPKHKKFKWKICVSPAKOLPGRIHFGSGYDNDPRGNLVYRFDVKDE 300
Qy 301 TYDLYQQCDAQPGASGSGVYVVMWKRQQQKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Db 301 TYDLYQQCDAQPGASGSGVYVVMWKRQQQKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Qy 361 ITPLKVAQICYWIKNYLDCREG 383
Db 361 ITPLKVAQICYWIKNYLDCREG 383

RESULT 11

US-09-072-384-15

; Sequence 15, Application US/09072384

; Patent No. 6153420

; GENERAL INFORMATION:

; APPLICANT: Sheppard, Paul O.

; TITLE OF INVENTION: SERINE PROTEASE POLYPEPTIDES

; AND MATERIALS AND METHODS FOR MAKING THEM

NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: ZymoGenetics, Inc.
STREET: 1201 Eastlake Avenue East
CITY: Seattle
STATE: WA
COUNTRY: USA
ZIP: 98102
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/072,384
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Parker, Gary E
REGISTRATION NUMBER: 31,648
REFERENCE/DOCKET NUMBER: 97-16C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-442-6673
TELEFAX: 206-442-6678
TELEX:

INFORMATION FOR SEQ ID NO: 15:

SEQUENCE CHARACTERISTICS:
LENGTH: 392 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
FRAGMENT TYPE: internal
FEATURE:
NAME/KEY: Signal Sequence
LOCATION: 1...19
OTHER INFORMATION:
US-09-072-384-15

Query Match 100.0%; Score 2080; DB 3; Length 392;

Best Local Similarity 100.0%; Pred. No. 5.6e-220;

Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY	1	MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
Db	1	MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
QY	61	VSSSCGPQCHKGTPPTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS	120
Db	61	VSSSCGPQCHKGTPPTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS	120
QY	121	SGSKRRKQIYGVDSRFSIFGKDFLLNYPSTSVKLSCTGCTGLVAEKHVLTAACHIDG	180
Db	121	SGSKRRKQIYGVDSRFSIFGKDFLLNYPSTSVKLSCTGCTGLVAEKHVLTAACHIDG	180
QY	181	KTVVKGTKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND	240
Db	181	KTVVKGTKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND	240
QY	241	IGMDYDVALLELKKPKHKKFMKIGVSPPAKQLPGGRIFHFGSYNDPGRNLVYRFDVKDE	300
Db	241	IGMDYDVALLELKKPKHKKFMKIGVSPPAKQLPGGRIFHFGSYNDPGRNLVYRFDVKDE	300
QY	301	TYDLLYQQCDAQPCASGSGVYVVMWKRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR	360
Db	301	TYDLLYQQCDAQPCASGSGVYVVMWKRQQQKWERKIIGIFSGHQWDMNGSPQDFNVAVR	360
QY	361	ITPLKQAQICYWIKNYLDREG 383	
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RESULT 12
US-09-072-384-2
Sequence 2, Application US/09072384
Patent No. 6153420
GENERAL INFORMATION:
APPLICANT: Sheppard, Paul O.
TITLE OF INVENTION: SERINE PROTEASE POLYPEPTIDES
AND MATERIALS AND METHODS FOR MAKING THEM
TITLE OF INVENTION: 18
NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: ZymoGenetics, Inc.
STREET: 1201 Eastlake Avenue East
CITY: Seattle
STATE: WA
COUNTRY: USA
ZIP: 98102
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/072,384
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Parker, Gary E
REGISTRATION NUMBER: 31,648
REFERENCE/DOCKET NUMBER: 97-16C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-442-6673
TELEFAX: 206-442-6678
TELEX:

INFORMATION FOR SEQ ID NO: 2:

SEQUENCE CHARACTERISTICS:
LENGTH: 392 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
FRAGMENT TYPE: internal
FEATURE:
NAME/KEY: Signal Sequence
LOCATION: 1...19
OTHER INFORMATION:
US-09-072-384-2

Query Match 98.3%; Score 2044; DB 3; Length 392;

Best Local Similarity 98.4%; Pred. No. 5.1e-216;

Matches 377; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY	1	MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
Db	1	MAGIPGLLFLFLLCAVGVSPYSAPWKTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE	60
QY	61	VSSSCGPQCHKGTPPTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS	120
Db	61	VSSSCGPQCHKGTPPTYEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS	120
QY	121	SGSKRRKQIYGVDSRFSIFGKDFLLNYPSTSVKLSCTGCTGLVAEKHVLTAACHIDG	180
Db	121	SGSKRRKQIYGVDSRFSIFGKDFLLNYPSTSVKLSCTGCTGLVAEKHVLTAACHIDG	180
QY	181	KTVVKGTKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND	240
Db	181	KTVVKGTKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKRTHVPKGIKGNAND	240
QY	241	IGMDYDVALLELKKPKHKKFMKIGVSPPAKQLPGGRIFHFGSYNDPGRNLVYRFDVKDE	300

REFERENCE/DOCKET NUMBER: 3396.214-US
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 212-867-0123
 TELEFAX: 212-867-0298
 INFORMATION FOR SEQ ID NO: 1:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 222 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 US-08-090-048-1

Query Match 8.5%; Score 177; DB 1; Length 222;
 Best Local Similarity 25.5%; Pred. No. 7.3e-11;
 Matches 60; Conservative 27; Mismatches 104; Indels 44; Gaps 9;

Qy	148	YFSTSVKLST---GCTGLVAEKHVLTAACHIDGKT--YVKGTKLRVGFLLKPKFKDGG	203
Db	17	YPYRAIVHSSIGSCTGMIGPKTVATAGHCIIYDTSSGSPAGTATVSPG-----	66
Qy	204	RGANDTSAMPEQMKFQWIRKTHVPKGIKGNANDIGMDYDVALLELKKPHKPKFKMI	263
Db	67	---RNGTS-----YPYGSVKSTRYFIPSGWRSGNTN-----YDYGAIELSEPIGNTVG YF	113
Qy	264	GVSPPAQLPGRIHFGSYDNDPCNLVYRCD--VKDETYDLLYQQCDAPGASGSGVY	321
Db	114	GYSTTSSLVGTVTTISGYPGDKTAGTQWQHSPIAISEYKIQYAM-DTYGGQSGSPVF	172
Qy	322	VRMWKROQKWKERII---GIFSGHQWDMNGSPQDFNVAVRITPLKYAQICW	372
Db	173	EQSSSRINCSPCSLAVHTNGVYGG-----SSYNRGTRITKEVFDNLTNW	217

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: July 1, 2005, 21:08:25 ; Search time 88.2836 Seconds
(without alignments)
1673.692 Million cell updates/sec

Title: US-09-658-677-18
Perfect score: 2080
Sequence: 1 MAGIPGLLFLFLCAVQ.....LKQAICYWIKGNVLDREG 383

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1722976 seqs, 385795295 residues
Total number of hits satisfying chosen parameters: 1722976

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications AA:*

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2:	/cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep.*
3:	/cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep.*
4:	/cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep.*
5:	/cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep.*
6:	/cgn2_6/ptodata/1/pubpaa/PCTUS_PUBCOMB.pep.*
7:	/cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep.*
8:	/cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep.*
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11:	/cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep.*
12:	/cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep.*
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17:	/cgn2_6/ptodata/1/pubpaa/US10E_PUBCOMB.pep.*
18:	/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep.*
19:	/cgn2_6/ptodata/1/pubpaa/US11A_PUBCOMB.pep.*
20:	/cgn2_6/ptodata/1/pubpaa/US11_NEW_PUB.pep.*
21:	/cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep.*
22:	/cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	2080	100.0	383	9	US-09-765-205-12
2	2080	100.0	383	9	US-09-909-320-261
3	2080	100.0	383	9	US-09-909-088B-261
4	2080	100.0	383	9	US-09-905-291A-261
5	2080	100.0	383	9	US-09-902-853-261
6	2080	100.0	383	9	US-09-907-824-261
7	2080	100.0	383	9	US-09-907-841-261
8	2080	100.0	383	10	US-09-904-011-261
9	2080	100.0	383	10	US-09-903-640-261
10	2080	100.0	383	10	US-09-908-093-261
11	2080	100.0	383	10	US-09-906-742-261

12	2080	100.0	383	10	US-09-906-838-261	Sequence 261, App
13	2080	100.0	383	10	US-09-907-613-261	Sequence 261, App
14	2080	100.0	383	10	US-09-907-942-261	Sequence 261, App
15	2080	100.0	383	10	US-09-904-859-261	Sequence 261, App
16	2080	100.0	383	10	US-09-909-204-261	Sequence 261, App
17	2080	100.0	383	10	US-09-904-820-261	Sequence 261, App
18	2080	100.0	383	10	US-09-904-786-261	Sequence 261, App
19	2080	100.0	383	10	US-09-906-646-261	Sequence 261, App
20	2080	100.0	383	10	US-09-906-700-261	Sequence 261, App
21	2080	100.0	383	10	US-09-903-788-261	Sequence 261, App
22	2080	100.0	383	10	US-09-902-903-261	Sequence 261, App
23	2080	100.0	383	10	US-09-903-749A-261	Sequence 261, App
24	2080	100.0	383	10	US-09-904-119-261	Sequence 261, App
25	2080	100.0	383	10	US-09-904-956-261	Sequence 261, App
26	2080	100.0	383	10	US-09-902-736-261	Sequence 261, App
27	2080	100.0	383	10	US-09-907-794-261	Sequence 261, App
28	2080	100.0	383	10	US-09-903-943-261	Sequence 261, App
29	2080	100.0	383	10	US-09-904-462-261	Sequence 261, App
30	2080	100.0	383	10	US-09-907-925-261	Sequence 261, App
31	2080	100.0	383	10	US-09-903-692-261	Sequence 261, App
32	2080	100.0	383	10	US-09-903-520-261	Sequence 261, App
33	2080	100.0	383	10	US-09-905-056-261	Sequence 261, App
34	2080	100.0	383	10	US-09-984-130-45	Sequence 45, Appl
35	2080	100.0	383	10	US-09-984-130-126	Sequence 126, App
36	2080	100.0	383	10	US-09-909-064-261	Sequence 261, App
37	2080	100.0	383	10	US-09-904-553-261	Sequence 261, App
38	2080	100.0	383	10	US-09-905-381-261	Sequence 261, App
39	2080	100.0	383	10	US-09-904-485-261	Sequence 261, App
40	2080	100.0	383	10	US-09-905-348-261	Sequence 261, App
41	2080	100.0	383	10	US-09-905-088-261	Sequence 261, App
42	2080	100.0	383	10	US-09-907-575-261	Sequence 261, App
43	2080	100.0	383	10	US-09-905-075-261	Sequence 261, App
44	2080	100.0	383	10	US-09-902-759-261	Sequence 261, App
45	2080	100.0	383	10	US-09-902-634-261	Sequence 261, App

ALIGNMENTS

RESULT 1
US-09-765-205-12
; Sequence 12, Application US/09765205
; Patent No. US20020034800A1
; GENERAL INFORMATION:
; APPLICANT: Cao, Li
; TITLE OF INVENTION: BONE MARROW SECRETED PROTEINS AND POLYNUCLEOTIDES
; FILE REFERENCE: 1458.004/200130.449
; CURRENT APPLICATION NUMBER: US/09/765,205
; CURRENT FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: US/09/212,440
; PRIOR FILING DATE: 1998-12-16
; NUMBER OF SEQ ID NOS: 46
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 12
; LENGTH: 383
; TYPE: PRT
; ORGANISM: human
; US-09-765-205-12

Query Match	100.0%;	Score	2080;	DB	9;	Length	383;
Best Local Similarity	100.0%;	Pred. No.	1.2e-194;				
Matches	383;	Conservative	0;	Mismatches	0;	Indels	0;
Qy	1	MAGIPGLLFLFLCAVGQVSPYSA	PWKPTWPAYRLPVVLPQSTLNLA	KPDFGA	EAKLE	60	
Db	1	MAGIPGLLFLFLCAVGQVSPYSA	PWKPTWPAYRLPVVLPQSTLNLA	KPDFGA	EAKLE	60	
Qy	61	VSSSCGPQCHKGTPLPTYEAKQYLS	YETLYANGSRTETQVGIIYILSSG	BGAQHRD	SGS	120	
Db	61	VSSSCGPQCHKGTPLPTYEAKQYLS	YETLYANGSRTETQVGIIYILSSG	BGAQHRD	SGS	120	
Qy	121	SGKSRKRQIYGVDSRPSIFGKDFLNL	PNPSTSVKLSKTGCTGTLVAE	KHVLTAAC	HIHDG	180	

Db 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
Qy 181 KTYVKGTKLRVGFLEKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTHPKGIKGNAND 240
Db 181 KTYVKGTKLRVGFLEKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTHPKGIKGNAND 240
Qy 241 IGMDYDVALLELKKPHKRKFMKIGVSPPAKQLPGGRIFHSGYDNDPRGNLVYRFDVKDE 300
Db 241 IGMDYDVALLELKKPHKRKFMKIGVSPPAKQLPGGRIFHSGYDNDPRGNLVYRFDVKDE 300
Qy 301 TYDLLYQCCDAQPCASGSGVYVVMWKRQOQKWERKIIIGIFSGHGWVDMNGSPQDENVAVR 360
Db 301 TYDLLYQCCDAQPCASGSGVYVVMWKRQOQKWERKIIIGIFSGHGWVDMNGSPQDENVAVR 360
Qy 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 2
US-09-909-320-261
; Sequence 261, Application US/09909320
; Patent No. US20020132240A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A. Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/909,320
; CURRENT FILING DATE: 2002-01-04
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214

; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-909-320-261

Query Match 100.0%; Score 2080; DB 9; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194; Indels 0; Gaps 0;
Matches 383; Conservative 0; Mismatches 0;

Qy 1 MAGIPGLLELLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAQPDGAEAKLE 60
Db 1 MAGIPGLLELLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAQPDGAEAKLE 60
Qy 61 VSSSCGPOCHKGTPLTYBEAKQYLSYETLYANGSRSTETQVGIYILSSSGDGAQHRDGS 120
Db 61 VSSSCGPOCHKGTPLTYBEAKQYLSYETLYANGSRSTETQVGIYILSSSGDGAQHRDGS 120
Qy 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
Db 121 SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLSCTGTGLVAEKHVLTAACHIDG 180
Qy 181 KTYVKGTKLRVGFLEKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTHPKGIKGNAND 240
Db 181 KTYVKGTKLRVGFLEKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTHPKGIKGNAND 240
Qy 241 IGMDYDVALLELKKPHKRKFMKIGVSPPAKQLPGGRIFHSGYDNDPRGNLVYRFDVKDE 300
Db 241 IGMDYDVALLELKKPHKRKFMKIGVSPPAKQLPGGRIFHSGYDNDPRGNLVYRFDVKDE 300
Qy 301 TYDLLYQCCDAQPCASGSGVYVVMWKRQOQKWERKIIIGIFSGHGWVDMNGSPQDENVAVR 360
Db 301 TYDLLYQCCDAQPCASGSGVYVVMWKRQOQKWERKIIIGIFSGHGWVDMNGSPQDENVAVR 360
Qy 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 3
US-09-909-088B-261
; Sequence 261, Application US/09909088B
; Patent No. US20020146709A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A. Paul J.
; APPLICANT: Godowski, Paul J.


```
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/909/088B
; CURRENT FILING DATE: 2001-07-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-909-088B-261

Best Local Similarity 100.0%; Score 2080; DB 9; Length 383;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLLLCAVGVSPYAPWKTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLLLCAVGVSPYAPWKTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Qy 61 VSSSCGPOCHKGTPLPYEAKQYLSVETLYANGSRTEQVGIYILSSGSGAQHRDGS 120
Db 61 VSSSCGPOCHKGTPLPYEAKQYLSVETLYANGSRTEQVGIYILSSGSGAQHRDGS 120
Qy 121 SGSRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLTVAEKHVLTAACHIDG 180
Db 121 SGSRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLTVAEKHVLTAACHIDG 180

; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,291A
; CURRENT FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29

; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,291A
; CURRENT FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
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; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29

; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,291A
; CURRENT FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
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; PRIOR APPLICATION NUMBER: US 60/146,222
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; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
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	PRIOR APPLICATION NUMBER:	PCT/US99/28313
	PRIOR FILING DATE: 1999-11-30	
	PRIOR APPLICATION NUMBER: PCT/US99/28564	
	PRIOR FILING DATE: 1999-12-02	
	PRIOR APPLICATION NUMBER: PCT/US99/28565	
	PRIOR FILING DATE: 1999-12-02	
	PRIOR APPLICATION NUMBER: PCT/US99/30095	
	PRIOR FILING DATE: 1999-12-16	
	PRIOR APPLICATION NUMBER: PCT/US99/30911	
	PRIOR FILING DATE: 1999-12-20	
	PRIOR APPLICATION NUMBER: PCT/US99/30999	
	PRIOR FILING DATE: 1999-12-20	
	PRIOR APPLICATION NUMBER: PCT/US00/00219	
	PRIOR FILING DATE: 2000-01-05	
	NUMBER OF SEQ ID NOS: 423	
	SEQ ID NO 261	
	LENGTH: 383	
	TYPE: PRT	
	ORGANISM: Homo Sapien	
	US-09-905-291A-261	
	Query Match	100.0%; Score 2080; DB 9; Length 383;
	Best Local Similarity	100.0%; Pred. No. 1.2e-194; Indels 0; Gaps 0;
	Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	
Qy	1	MAGIPGLLFLFFLLCAVGVSPYSAPWKPTWPAYRLPVLPQSTLNLAQDFGAEAKLE 60
Db	1	MAGIPGLLFLFFLLCAVGVSPYSAPWKPTWPAYRLPVLPQSTLNLAQDFGAEAKLE 60
Qy	61	VSSCGPQCHKGTPLTPEEAKQVLSYETLVANGSRRTQVGIYIILSSSGDAQHRDGS 120
Db	61	VSSCGPQCHKGTPLTPEEAKQVLSYETLVANGSRRTQVGIYIILSSSGDAQHRDGS 120
Qy	121	SGKRRRQIYGYDSRSIFCKDFLLNYPFSTSVKLSGCTGTLVAERHVLTAACHIDG 180
Db	121	SGKRRRQIYGYDSRSIFCKDFLLNYPFSTSVKLSGCTGTLVAERHVLTAACHIDG 180
Qy	181	KTYVKGTKLRVGLFKPKFKDGGRGANDSTAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
Db	181	KTYVKGTKLRVGLFKPKFKDGGRGANDSTAMPEQMKFQWIRVKRTHVPKGIKGNAND 240
Qy	241	IGMDYVALLLEKPKHKKFKMKIIGVSPPAKQLPGRRIHFSGYDNDRPGNLVYRFCDVKDE 300
Db	241	IGMDYVALLLEKPKHKKFKMKIIGVSPPAKQLPGRRIHFSGYDNDRPGNLVYRFCDVKDE 300
Qy	301	TYDLLYQCCDAQPGASGSGVYVRWKKRQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
Db	301	TYDLLYQCCDAQPGASGSGVYVRWKKRQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
Qy	361	ITPLKYAQICWIKGNVLDRCRG 383
Db	361	ITPLKYAQICWIKGNVLDRCRG 383
RESULT 5		
US-09-902-853-261		
Sequence 261, Application US/09902853		
Publication No. US20020192659A1		
GENERAL INFORMATION:		
APPLICANT: Genentech, Inc.		
APPLICANT: Ashkenazi, Avi		
APPLICANT: Botstein, David		
APPLICANT: Desnoyers, Luc		
APPLICANT: Eaton, Dan L.		
APPLICANT: Ferrara, Napoleone		
APPLICANT: Filvaroff, Ellen		
APPLICANT: Fong, Sherman		
APPLICANT: Gao, Wei-Qiang		
APPLICANT: Gerber, Hanspeter		
APPLICANT: Gerritsen, Mary E.		
APPLICANT: Goddard, A.		
APPLICANT: Godowski, Paul J.		
APPLICANT: Grimaldi, Christopher J.		

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Qy 181 KTYVKGTKLRVGLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
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Db 181 KTYVKGTKLRVGLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
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Qy 241 IGMVDYALLELKKPHKFKMKIGVSPAKQLPGRIHFSGYNDNRPGNLVYRFDVKDE 300
      |||||
Db 241 IGMVDYALLELKKPHKFKMKIGVSPAKQLPGRIHFSGYNDNRPGNLVYRFDVKDE 300
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Qy 301 TYDLLYQCCDAQPGASGGVYVVMWKQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
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Db 301 TYDLLYQCCDAQPGASGGVYVVMWKQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
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Qy 361 ITPLKYAQCWIKGNLYDCREG 383
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Db 361 ITPLKYAQCWIKGNLYDCREG 383
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RESULT 6
US-09-907-824-261
; Sequence 261, Application US/09907824
; Publication No. US20020197671A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Hurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,824
; CURRENT FILING DATE: 2001-07-17
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
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; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-907-824-261

Query Match 100.0%; Score 2080; DB 9; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLPFLLCAGVQSPYSAPWKPWTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
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Db 1 MAGIPGLLFLPFLLCAGVQSPYSAPWKPWTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
      |||||
Qy 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDSGS 120
      |||||
Db 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDSGS 120
      |||||
Qy 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTGLVAEKHVLTAHCIIHDG 180
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Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSCTGTGLVAEKHVLTAHCIIHDG 180
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Qy 181 KTYVKGTKLRVGLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
      |||||
Db 181 KTYVKGTKLRVGLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTVPKGIKGNAND 240
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Qy 361 ITPLKYAQCWIKGNLYDCREG 383
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Db 361 ITPLKYAQCWIKGNLYDCREG 383
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RESULT 7
US-09-907-841-261
; Sequence 261, Application US/09907841
; Publication No. US20020198366A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
```

```

; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,841
; CURRENT FILING DATE: 2001-11-20
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
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; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-907-841-261

Query Match          100.0%; Score 2080; DB 9; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1  MAGIPGLLFLFLCAVGQVSPYSAPWKPPTWPAYRLPVVLPQSTLNLAQPDGAEAKLE 60
      |||
Db      1  MAGIPGLLFLFLCAVGQVSPYSAPWKPPTWPAYRLPVVLPQSTLNLAQPDGAEAKLE 60
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Qy      61  VSSSCGPQCHGTPLTYEBAKOYLSYETUYANGSRTEQVGIIYILSSSGDGAQHRDGS 120
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Db      61  VSSSCGPQCHGTPLTYEBAKOYLSYETUYANGSRTEQVGIIYILSSSGDGAQHRDGS 120
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Qy      121  SGKSRKRQIYGYDYSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEKHVLTAACHIDG 180
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Db      121  SGKSRKRQIYGYDYSRFSIFGKDFLLNYPFSTSVKLSCTGCTGLVAEKHVLTAACHIDG 180
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Qy      181  KTVYVGTQKLRVGLFKPKFKDGGGRGANDSTSAMPEQMKFQWIRVVRKTRHVPKGIKGNAND 240
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Db      181  KTVYVGTQKLRVGLFKPKFKDGGGRGANDSTSAMPEQMKFQWIRVVRKTRHVPKGIKGNAND 240
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Qy      241  IGMDDYALLELKKPHKPKMKIYGVSPAPKQLPGGRIHFSGYNDPRGNLYVRFCDVKDE 300
      |||
Db      241  IGMDDYALLELKKPHKPKMKIYGVSPAPKQLPGGRIHFSGYNDPRGNLYVRFCDVKDE 300
      |||

Qy      301  TYDLLYQCDQAQPGASGSGVYVRWKRQQQKWERKLIIGIFSGHQWDMNGSPQDFNVAVR 360
      |||
Db      301  TYDLLYQCDQAQPGASGSGVYVRWKRQQQKWERKLIIGIFSGHQWDMNGSPQDFNVAVR 360
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Qy      361  ITPLKYAQICYWIKGNYLDCREG 383
      |||
Db      361  ITPLKYAQICYWIKGNYLDCREG 383
      |||

RESULT 8
US-09-904-011-261
; Sequence 261, Application US/09904011
; Publication No. US20030003530A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/904,011
; CURRENT FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
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PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-904-011-261

Query Match 100.0%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLLFLLCAVGVSPYSAPWKPPTWYRLPVVLPSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLLFLLCAVGVSPYSAPWKPPTWYRLPVVLPSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDGS 120
DB 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDGS 120

QY 121 SGKRRKROIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
DB 121 SGKRRKROIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG 180

QY 181 KTVVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240
DB 181 KTVVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240

QY 241 IGMVDYALLEKPKHKKFKEMKIGVSPPAKQLPGGRIFHSGYDNDPRGNLVYRFDVKDE 300
DB 241 IGMVDYALLEKPKHKKFKEMKIGVSPPAKQLPGGRIFHSGYDNDPRGNLVYRFDVKDE 300

QY 301 TYDLLYQQCDAQPGASGSGVYVVMKRRQOKWERRKIIGIFSGHQWDMNGSPQDFNVAVR 360
DB 301 TYDLLYQQCDAQPGASGSGVYVVMKRRQOKWERRKIIGIFSGHQWDMNGSPQDFNVAVR 360

RESULT 9

US-09-903-640-261
Sequence 261, Application US/09903640
Publication No. US20030017463A1
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Pao, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.

TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/903,640
CURRENT FILING DATE: 2001-07-11
PRIOR APPLICATION NUMBER: 09/665,350
PRIOR FILING DATE: 2000-09-18
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-903-640-261

Query Match 100.0%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLLFLLCAVGVSPYSAPWKPPTWYRLPVVLPSTLNLAKEPFGAEAKLE 60
DB 1 MAGIPGLLLFLLCAVGVSPYSAPWKPPTWYRLPVVLPSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDGS 120
DB 61 VSSSCGPOCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDGS 120

QY 121 SGKRRKROIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
DB 121 SGKRRKROIYGYDSRFSIFGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG 180

QY 181 KTVVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240
DB 181 KTVVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKFQWIRVKTTHVPKGIKGNAND 240

QY 241 IGMVDYALLEKPKHKKFKEMKIGVSPPAKQLPGGRIFHSGYDNDPRGNLVYRFDVKDE 300
DB 241 IGMVDYALLEKPKHKKFKEMKIGVSPPAKQLPGGRIFHSGYDNDPRGNLVYRFDVKDE 300

QY 301 TYDLLYQQCDAQPGASGSGVYVVMKRRQOKWERRKIIGIFSGHQWDMNGSPQDFNVAVR 360
DB 301 TYDLLYQQCDAQPGASGSGVYVVMKRRQOKWERRKIIGIFSGHQWDMNGSPQDFNVAVR 360

QY 361 ITPKYAQICYWIKGNLYDCREG 383
DB 361 ITPKYAQICYWIKGNLYDCREG 383

RESULT 10

US-09-908-093-261
Sequence 261, Application US/09908093
Publication No. US20030017498A1
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Pao, Nicholas F.
APPLICANT: Roy, Margaret Ann

APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/908,093
CURRENT FILING DATE: 2001-07-17
PRIOR APPLICATION NUMBER: 09/665,350
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/30095
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 261
LENGTH: 383
TYPE: PRT
ORGANISM: Homo Sapien
US-09-908-093-261

Query Match 100.0%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAGIPGLLFLFLLCAVGQVSPYAPWKPTPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGQVSPYAPWKPTPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

Qy 61 VSSSCGPQCHKGTPLPYERAKQVLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS 120
Db 61 VSSSCGPQCHKGTPLPYERAKQVLSYETLYANGSRTEQVGIYILSSSGDGAQHRDGS 120

Qy 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSGTCTGLTVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGVDSRFSIFGKDFLLNYPSTSVKLSGTCTGLTVAEKHVLTAACHIDG 180

Qy 181 KTVVKGQKLRVGLFKPKFGDGRGANDTSAMPEQMKFQIRVKRTHVPKGIKGNAND 240
Db 181 KTVVKGQKLRVGLFKPKFGDGRGANDTSAMPEQMKFQIRVKRTHVPKGIKGNAND 240

Qy 241 IGMDDYALLELKKPKRKFPMKIGVSPPAKQLPGRIHFSGYDNDRPNLVRFCVDKDE 300

Db 241 IGMDDYALLELKKPKRKFPMKIGVSPPAKQLPGRIHFSGYDNDRPNLVRFCVDKDE 300
Qy 301 TYDLLYQQCDAQPGASGSGVYVMMKROQKWKRIIGIFSGHQWYDMNGSPQDFNVAVR 360
Db 301 TYDLLYQQCDAQPGASGSGVYVMMKROQKWKRIIGIFSGHQWYDMNGSPQDFNVAVR 360
Qy 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 11
US-09-906-742-261
Sequence 261, Application US/09906742
Publication No. US20030023054A1
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Faoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/906,742
CURRENT FILING DATE: 2001-07-16
PRIOR APPLICATION NUMBER: 09/665,350
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02

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; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-906-742-261

Query Match      100.0%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPOCHKGTPPTYEEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDGS 120
Db 61 VSSSCGPOCHKGTPPTYEEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDGS 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
QY 181 KTVVGTQKLRVGFLLKPKFKDGGGRANDSTSAMPEQMKFQWIRVGRTHVPKGIKNAND 240
Db 181 KTVVGTQKLRVGFLLKPKFKDGGGRANDSTSAMPEQMKFQWIRVGRTHVPKGIKNAND 240
QY 241 IGWDYDYLLELKKPKFKEMKIGVSPPAKOLPGGRHSGYNDPRGNLVYFCDVKDE 300
Db 241 IGWDYDYLLELKKPKFKEMKIGVSPPAKOLPGGRHSGYNDPRGNLVYFCDVKDE 300
QY 301 TYDLLYQCCDAOPGASGSGYVVMWKRQOQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
Db 301 TYDLLYQCCDAOPGASGSGYVVMWKRQOQKWKRIIGIFSGHQWDMNGSPQDFNVAVR 360
QY 361 ITPKVAQICYTWIKGNLYDCREG 383
Db 361 ITPKVAQICYTWIKGNLYDCREG 383

RESULT 12
US-09-906-838-261
; Sequence 261, Application US/09906838
; Publication No. US20030027143A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
```

```
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/906,838
; CURRENT FILING DATE: 2001-07-16
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-906-838-261

Query Match      100.0%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
QY 61 VSSSCGPOCHKGTPPTYEEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDGS 120
Db 61 VSSSCGPOCHKGTPPTYEEAKQYLSYETLYANGSRTEQVGIYIILSSSGDGAQHRDGS 120
QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
QY 181 KTVVGTQKLRVGFLLKPKFKDGGGRANDSTSAMPEQMKFQWIRVGRTHVPKGIKNAND 240
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Db 181 KTVVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKQWIRVKRTHVPKGIKGNAND 240
QY 241 IGMDDYALLENLKKPHKRFKMGKIGVSPPAKQLPFGRIHFGSYDNDPRGNLVYRFDVKDE 300
Db 241 IGMDDYALLENLKKPHKRFKMGKIGVSPPAKQLPFGRIHFGSYDNDPRGNLVYRFDVKDE 300
QY 301 TYDLLYQOCDAQPGASGSGVYVYRMKROQOKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Db 301 TYDLLYQOCDAQPGASGSGVYVYRMKROQOKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
QY 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 13

US-09-907-613-261
; Sequence 261, Application US/09907613
; Publication No. US20030027145A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,613
; CURRENT FILING DATE: 2001-07-17
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564

; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-907-613-261

Query Match 100.0%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPGLLFLFLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60

QY 61 VSSSCGPOCHKGTPLTYEEAKQYLSYETLYANGSRTEQVGIYIILSSGDSGAQHRDSGS 120
Db 61 VSSSCGPOCHKGTPLTYEEAKQYLSYETLYANGSRTEQVGIYIILSSGDSGAQHRDSGS 120

QY 121 SGKSRKRQIYGDSRFSIFGKDFLLNYPFSTSVKLSCTCTGTTLVAEKHVLTAACHIDG 180
Db 121 SGKSRKRQIYGDSRFSIFGKDFLLNYPFSTSVKLSCTCTGTTLVAEKHVLTAACHIDG 180

QY 181 KTVVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKQWIRVKRTHVPKGIKGNAND 240
Db 181 KTVVGTQKLRVGLFKPKFDGGRGANDSTSAMPEQMKQWIRVKRTHVPKGIKGNAND 240

QY 241 IGMDDYALLENLKKPHKRFKMGKIGVSPPAKQLPFGRIHFGSYDNDPRGNLVYRFDVKDE 300
Db 241 IGMDDYALLENLKKPHKRFKMGKIGVSPPAKQLPFGRIHFGSYDNDPRGNLVYRFDVKDE 300

QY 301 TYDLLYQOCDAQPGASGSGVYVYRMKROQOKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360
Db 301 TYDLLYQOCDAQPGASGSGVYVYRMKROQOKWERKIIGIFSGHQWYDMNGSPQDFNVAVR 360

QY 361 ITPLKYAQICYWIKGNLYDCREG 383
Db 361 ITPLKYAQICYWIKGNLYDCREG 383

RESULT 14

US-09-907-942-261
; Sequence 261, Application US/09907942
; Publication No. US20030027146A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.


```

; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,942
; CURRENT FILING DATE: 2002-01-22
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
; US-09-907-942-261

Query Match 100.0%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194; Indels 0; Gaps 0;
Matches 383; Conservative 0; Mismatches 0;

QY 1 MAGIPGLFLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAQKPDFAEAKLE 60
DB 1 MAGIPGLFLFLFLLCAVGVSPYSAPWKPWPAYRLPVVLPQSTLNLAQKPDFAEAKLE 60
QY 61 VSSSCGPGCHGTPLPTPYEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDSSG 120
DB 61 VSSSCGPGCHGTPLPTPYEAKQYLSYETLYANGSRTETQVGIYILSSSGDGAQHRDSSG 120
QY 121 SGKSRKRIQYGVDSRFSGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
DB 121 SGKSRKRIQYGVDSRFSGKDFLLNYPSTSVKLSGTCTGLVAEKHVLTAACHIDG 180
QY 181 KTVVKGQTKLRVGLFKPKDKGGRGANDSTSAMPEQMKFQWIRVKKTHVPKGIKNAND 240
DB 181 KTVVKGQTKLRVGLFKPKDKGGRGANDSTSAMPEQMKFQWIRVKKTHVPKGIKNAND 240

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QY 241 IGMDYDVALLELKKPKRKMKGVSPPAKQLPGGRHFGSYDNDPRGNLVYRFDVKDE 300
DB 241 IGMDYDVALLELKKPKRKMKGVSPPAKQLPGGRHFGSYDNDPRGNLVYRFDVKDE 300
QY 301 TYDLLYQQCDAOPGASGSGVYVVMWKRQOQKWKRIIGIFSGHQWVDMNGSPQDFNVAVR 360
DB 301 TYDLLYQQCDAOPGASGSGVYVVMWKRQOQKWKRIIGIFSGHQWVDMNGSPQDFNVAVR 360
QY 361 ITPKTYAQICYWIKGNLYDCREG 383
DB 361 ITPKTYAQICYWIKGNLYDCREG 383

RESULT 15
US-09-904-859-261
; Sequence 261, Application US/09904859
; Publication No. US200300306060A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/904,859
; CURRENT FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30

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; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 261
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-904-859-261

Query Match      100.0%; Score 2080; DB 10; Length 383;
Best Local Similarity 100.0%; Pred. No. 1.2e-194;
Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1  MAGIPGLLFLFLLCAVGVSPYSAPKPTWPAYRLPVVLPOSTLNLAKEPFGAEAKLE 60
Db      1  MAGIPGLLFLFLLCAVGVSPYSAPKPTWPAYRLPVVLPOSTLNLAKEPFGAEAKLE 60

Qy     61  VSSSCGPGCHKGTPPTYEAKQVLSYETLYANGSRRTETQVGIYILSSSGDGAQHRDGS 120
Db     61  VSSSCGPGCHKGTPPTYEAKQVLSYETLYANGSRRTETQVGIYILSSSGDGAQHRDGS 120

Qy    121  SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLTGCTGTIVAEPKHVLTAAHCIHDG 180
Db    121  SGKRRKRQIYGVDSRFSIFGKDFLLNYPFSTSVKLTGCTGTIVAEPKHVLTAAHCIHDG 180

Qy    181  KTYVKGTKLRLVGLKPKFKDGGGANDSTSAMPEQMKFQWIRVKTHTVPEKGIKGNAND 240
Db    181  KTYVKGTKLRLVGLKPKFKDGGGANDSTSAMPEQMKFQWIRVKTHTVPEKGIKGNAND 240

Qy    241  IGMDDYDYLLELKKPKRKPWKIGVSPPAKQLPGRIHFSGYDNDRPGNLVYRFCVVKDE 300
Db    241  IGMDDYDYLLELKKPKRKPWKIGVSPPAKQLPGRIHFSGYDNDRPGNLVYRFCVVKDE 300

Qy    301  TYDLLYQQCDAQPGASGSGVYVRMWKQQQKWERKIIGIFSGHQWVDMNGSPQDFNVAVR 360
Db    301  TYDLLYQQCDAQPGASGSGVYVRMWKQQQKWERKIIGIFSGHQWVDMNGSPQDFNVAVR 360

Qy    361  ITPLKYAQICYWKGNVLDREG 383
Db    361  ITPLKYAQICYWKGNVLDREG 383
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Search completed: July 1, 2005, 21:32:18
Job time : 89.2836 secs

GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: July 1, 2005, 20:54:23 ; Search time 20.3479 Seconds
(without alignments)
1811.048 Million cell updates/sec

Title: US-09-658-677-18

Perfect score: 2080

Sequence: 1 MAGIPGLFLFLFLCAVGQ.....LKQAICYWIKGNLYLDCREG 383

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

PIR 79:.*
1: pir1.*
2: pir2.*
3: pir3.*
4: pir4.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	194	9.3	316	2 A45134	endopeptidase (EC
2	135.5	6.5	313	2 A35122	metalloproteinase
3	131	6.3	218	2 E97915	choline binding pr
4	120.5	5.8	482	1 EXRT	coagulation factor
5	119	5.7	269	2 A26823	pancreatic elastas
6	117	5.6	522	2 T29767	hypothetical prote
7	115.5	5.6	289	2 B26823	pancreatic elastas
8	115.5	5.6	271	2 A25528	pancreatic elastas
9	115	5.5	258	4 S70439	pancreatic elastas
10	115	5.5	267	4 A56615	probable pancreati
11	114.5	5.5	238	1 TRW5Y	trypsin-like prote
12	114.5	5.5	285	2 C95045	choline binding pr
13	114	5.5	246	1 DBHU	complement factor
14	113	5.4	266	1 ELPG	pancreatic elastas
15	113	5.4	266	1 ELRT1	pancreatic elastas
16	112.5	5.4	259	1 TRSMG	trypsin (EC 3.4.21
17	112.5	5.4	273	2 E85765	hypothetical prote
18	112.5	5.4	273	2 H64915	putative protease
19	111.5	5.4	488	1 EXHU	coagulation factor
20	111	5.3	761	2 JC5759	brain-specific ser
21	110	5.3	269	2 C26823	pancreatic elastas
22	110	5.3	492	1 EXBO	coagulation factor
23	110	5.3	1582	2 T15308	hypothetical prote
24	109.5	5.3	405	2 T35117	probable secreted
25	109	5.2	278	2 AH0282	probable peptidas
26	108	5.2	583	2 A29154	complement factor
27	108	5.2	786	1 A47547	serine proteinase
28	108	5.2	1047	2 A55617	masquerade precurs
29	107.5	5.2	236	2 A28566	T-cell suppressor

30 107.5 5.2 686 1 A59271 Ra-reactive factor
31 105 5.0 274 2 S40004 trypsin-related pr
32 103.5 5.0 271 1 ELRT2 pancreatic elastas
33 103.5 5.0 416 1 KFBQ coagulation factor
34 102.5 4.9 1238 2 T34929 hypothetical prote
35 100 4.8 272 2 JC4170 trypsin-like prote
36 98.5 4.7 267 2 S40006 trypsin (EC 3.4.21
37 98 4.7 548 2 D82175 probable trypsin V
38 98 4.7 624 2 T02289 probable polygalac
39 98 4.7 855 2 JC7731 membrane-bound arg
40 97.5 4.7 409 2 T35118 probable secreted
41 97.5 4.7 452 1 A30351 coagulation factor
42 97.5 4.7 747 2 I51579 complement factor
43 97 4.7 1019 2 A38738 coagulation factor
44 96.5 4.6 259 2 S68424 allergen Der f III
45 96.5 4.6 275 2 I46712 factor IX - rabbit

ALIGNMENTS

RESULT 1
A45134
endopeptidase (EC 3.4.-.-), glutamate-specific - Bacillus licheniformis
C:Species: Bacillus licheniformis
C:Date: 10-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 16-Aug-2004
C:Accession: A45134, S23078
R:Kakudo, S.; Kikuchi, N.; Kitadokoro, K.; Fujiwara, T.; Nakamura, E.; Okamoto, H.; Shin, J. Biol. Chem. 267, 23782-23788, 1992
A:Title: Purification, characterization, cloning, and expression of a glutamic acid-speci
A:Reference number: A45134; MUID:93054737; PMID:1429718
A:Accession: A45134
A>Status: preliminary
A:Molecule type: DNA
A:Residues: 1-316 <KAK>
A:Cross-references: UNIPROT:P80057; GB:D10060; NID:g216263; PIDN:BAA00949.1; PID:d100141;
A:Experimental source: ATCC 14580
A:Note: sequence extracted from NCBI backbone (NCBIN:118784, NCBIP:118785)
R: Svendsen, I.; Breddam, K.
Eur. J. Biochem. 204, 165-171, 1992
A:Title: Isolation and amino acid sequence of a glutamic acid specific endopeptidase from
A:Reference number: S23078; MUID:92155199; PMID:1346764
A:Accession: S23078
A>Status: preliminary
A:Molecule type: protein
A:Residues: 95-316 <SVE>
C:Superfamily: Glutamyl endopeptidase, V8 type
C:Keywords: hydrolase

Query Match 9.3%; Score 194; DB 2; Length 316;
Best Local Similarity 24.0%; Pred. NO. 1.7e-08;
Matches 79; Conservative 43; Mismatches 143; Indels 64; Gaps 15;
QY 63 SSGCPCHKGTPL---PTVEAKQVLSYETLVANGSRTEQTGVGIYILSSSGCAQHRDSG 119
Db 28 AQAPSPH--TPVSSDPSY-KAETSVTYP-----NIKSDQVGLYSKAFGTGKVNETKE 79
QY 120 SSGKSRKRQIYGDSRSPFPGKDFLLN-----YPFSTSVKLST---GCTGLTVAEKHV 170
Db 80 KAEKSPAKAPY--SIKSVIGSDDRTRVNTTAYPYRAIVHSSISGSGCTGMIGPKTV 136
QY 171 LTAACIHGCKT-YVKGTKLRVGLFKPKFGKGRANDSTSMPEQMKFQWTRVKRTHV 229
Db 137 ATAGHCIDYTSSGSGFAGTATVSPG-----RNGTS-----YPYGSVKSTRFYI 178
QY 230 PKGWTIGNANDIGMDYDVALLEKPKRKFMIKIGVSPPAKQLPGGRIHFSGYDNDRPGN 289
Db 179 PGWRSNGTN-----YDYGAIELSEPIGNVTGVFGYSYTTSSLVGTVTIVISPGDKTAG 233
QY 290 LVYRFCD--VKDETLLYQQCDAQPFASGSGGVYVRMKRQQQKWERKI---GIFSGH 343
Db 234 TQWHSGPATSETYKLYQAM-DTYGQSGSPVFEQSSSRSTNCSCGPCSLAVHTNGVYGG- 291

QY 344 QWDMNGSPQDFNVAVRITPLKVAQICYW 372
Db 292 -----SSYNRGRTITKEVFDNLTNW 311

RESULT 2
A:Title: metalloproteinase (EC 3.4.4.-) mpr precursor, extracellular - Bacillus subtilis
C:Species: Bacillus subtilis
C:Date: 27-Jul-1990 #sequence revision 27-Jul-1990 #text_change 16-Aug-2004
C:Accession: A35122; I40010; A69660
R:Sloma, A.; Rudolph, C.F.; Rufo Jr., G.A.; Sullivan, B.J.; Theriault, K.A.; Ally, D.; J. Bacteriol. 172, 1024-1029, 1990
A:Title: Gene encoding a novel extracellular metalloprotease in Bacillus subtilis.
A:Reference number: A35122; MUID:90130256; PMID:2105291
A:Accession: A35122
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-313 <SLO>
A:Cross-references: UNIPROT:P39790; GB:L10505; NID:G143209; PIDN:AAA22604.1; PID:G143210
R:Smith, H.; de Jong, A.; Bron, S.; Venema, G.
Gene 70, 351-361, 1988
A:Title: Characterization of signal-sequence-coding regions selected from the Bacillus subtilis genome.
A:Reference number: I39994; MUID:99108019; PMID:3145906
A:Accession: I40010
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-60, 65, 'L', '67', 'S', '69', 'AOK' <RES>
A:Cross-references: GB:M22916; NID:G143701; PIDN:AAA22832.1; PID:G143702
R:Kunst, P.; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Berton
C.; Bron, S.; Brouillet, S.; Bruchin, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Ch
A.; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
Nature 390, 249-256, 1997
A:Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Gallen
iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holeappel, S.; Hosono, S.; Hullo, M.F.
Koertter, P.; Koningsstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois,
A:Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maueel
Y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetelle
Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon,
A:Authors: Schleich, S.; Schroeter, R.; Scoffone, P.; Sekiguchi, J.; Sekowska, A.; Seron
akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, J.; Tosiato, V.; Uchiyama
T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, K
A:Authors: Yoshikawa, H.F.; Zumbstein, E.; Yoshikawa, H.; Danchin, A.
A:Title: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis.
A:Reference number: A69580; MUID:98044033; PMID:9384377
A:Accession: A69560
A:Status: preliminary; nucleic acid sequence not shown; translation not shown
A:Molecule type: DNA
A:Residues: 1-313 <KUN>
A:Cross-references: GB:Z99105; GB:AL009126; NID:G2632457; PIDN:CAB12018.1; PID:G1182176;
A:Experimental source: strain 168
C:Genetics:
A:Gene: mpr
C:Superfamily: Glutamyl endopeptidase, V8 type
C:Keywords: hydrolase

Query Match 6.5%; Score 135.5; DB 2; Length 313;
Best Local Similarity 22.2%; Pred. No. 0.0013;
Matches 77; Conservative 38; Mismatches 121; Indels 111; Gaps 17;

QY 72 GTPLPYVEAKQYLVETLYANGSRTEQVGIYLVSSGD--GAQHRDSSGSSKSRKQ 129
Db 29 GVPKAAENPQTSVNSGKEADATKNQT-----SKADQVAPYEGTGKTSK----- 75

QY 130 IYGVDSRF-----STFGKD-----FLNLYPSTSVKLSL-----GCTGTL 164
Db 76 LYGGQTELEKNIQLOPSSIIGTDERTRISSTSFPRATVQUSIKYKPNSTSYGCTGFL 135

QY 165 VAERKHVLTAAHCIIH-------DGKTYVKGTOKLRVGLKPKFKDGGRGAND 208
Db 136 VNPNTVVTAGHCYVSQDHGWASTTAAPGRNGSSYPGVY----- 175

QY 209 STSAMPEQMKFQIRVKRTHVPGWIKGN---ANDTGMVDYALLELKKPKHKKFMKIGV 265

Db 176 -SGTFYSVK-GWTBSKDTNYDYGAIKLNGSPGNTVGM-YGYRTNSSSP-----VGL 225
QY 266 SPPAKQLPGGRIHFSGYNDRPGNLVYRFDCKVDEYDLYLQQCDAQPGASGGVYVRMW 325
Db 226 SSSVTGFPCKDTFGTMMWSDTKPIR-----SAETYKLTYYTDTYGCQSGSPVY---- 272

QY 326 KRQOQKWERKIIFPSGHQWDMNGSPQDFNVAVRITPLKVAQICYW 372
Db 273 -RNYSDTGQTATAIHT-----NGG-SSYNLGTRVNDVFNNIQYW 310

RESULT 3
E97915
A:Title: choline binding protein G, truncation [imported] - Streptococcus pneumoniae (strain R6)
C:Species: Streptococcus pneumoniae
C:Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 09-Jul-2004
C:Accession: E97915
R:Hoskins, J.A.; Alborn Jr., W.; Arnold, J.; Blaszcak, L.; Burgett, S.; DeHoff, B.S.; E
e, R.; LeBlanc, D.J.; Lee, L.N.; Lefkowitz, E.J.; Lu, J.; Matsushima, P.; McAhren, S.; M
Y, P.; Sun, P.M.; Winkler, M.E.
J. Bacteriol. 183, 5709-5717, 2001
A:Authors: Yang, Y.; Young-Bellido, M.; Zhao, G.; Zook, C.; Baltz, R.H.; Jaskunas, S.R.;
A:Title: Genome of the Bacterium Streptococcus pneumoniae Strain R6.
A:Reference number: A97872; MUID:21429245; PMID:11544234
A:Accession: E97915
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-218 <KUR>
A:Cross-references: UNIPROT:Q8DR41; GB:AE007317; PIDN:AAK99153.1; PID:G15457907; GSPDB:G
C:Genetics:
A:Gene: cbpG-truncation

Query Match 6.3%; Score 131; DB 2; Length 218;
Best Local Similarity 24.2%; Pred. No. 0.0019;
Matches 59; Conservative 30; Mismatches 87; Indels 68; Gaps 10;

QY 143 DFLNYPSTSVKLSGCTG-----TLVAEKHVLTAACHIDGKTYVKGTOK 189
Db 5 DNTLQVPYSTSAWLSSKYVGADGMNVEGRGSANFKONLVITAHH-----NYR----- 54

QY 190 LRVGFLKPKFKDGGRGAND---STSAMPEQMKFQIRVKRTHVPGWIKGNANDIGMDYD 246
Db 55 -----HDYGEADDIYVLPVAVSPSQELFKIKVKEVRYLKEFRNLNSKD-AREYD 103

QY 247 YALLELKKPKHKKFKMGIVSPPAKQLPGGRIHFSGYNDRPGNLVYR-----CDVKDE 300
Db 104 LALLILEKPIGAKLTGLGLPTSQKNLTGITVITGYPS-----YNFKIHQWYTDKKQV 156

QY 301 TYD---LLYQOCDAQPGASGGVYVRMWKROQKWERKIIFPSGHQWDMNGSPQDFNV 357
Db 157 LSDDGMFLDYQVDTLEGSSGSTVY-----DASHRVVGVT-----LGDGANQINS 201

QY 358 AVRI 361
Db 202 AVKL 205

RESULT 4
EXPT
A:Title: cosulation factor Xa (EC 3.4.21.6) precursor - rat
C:Species: Rattus norvegicus (Norway rat)
C:Date: 31-Jan-1995 #sequence_revision 07-Feb-1997 #text_change 09-Jul-2004
C:Accession: S49075; J04670; PS0191; PS0190; I62745
R:Stanton, C.; Ross, P.; Hutson, S.; Wallin, R.
Thromb. Res. 80, 63-73, 1995
A:Title: Evidence for competition between vitamin K-dependent clotting factors for intrac
A:Reference number: A58498; MUID:96093366; PMID:8578539
A:Accession: S49075
A:Molecule type: mRNA
A:Residues: 1-482 <STAL>
A:Cross-references: UNIPROT:Q63207; EMBL:X79807; NID:G506600; PIDN:CAA56202.1; PID:G50660
A:Note: submitted to the EMBL Data Library, June 1994

```

R;Kawashima, I.; Tani, T.; Shimoda, K.; Takiguchi, Y.
DNA 6, 163-172, 1987
A;Title: Characterization of pancreatic elastase II cDNAs: two elastase II mRNAs are expressed from a single gene in rat pancreas
A;Reference number: A90958; MUID:87217962; PMID:3646943
A;Accession: A26823
A;Molecule type: mRNA
A;Residues: 1-269 <KW>
A;Cross-references: UNIPROT:P08419; GB:M16651; NID:g164441; PIDN:AAA31027.1; PID:g164442
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; serine proteinase
F;1-16/Domain: signal sequence #status predicted <SIG>
F;17-28/Domain: propeptide #status predicted <PRO>
F;29-269/Product: elastase II #status predicted <MAT>
F;73-262/Domain: trypsin homology <TRY>
F;73,121,216/Active site: His, Asp, Ser #status predicted

Query Match          5.7%; Score 119; DB 2; Length 269;
Best Local Similarity 27.5%; Pred No. 0.024;
Matches 60; Conservative 28; Mismatches 64; Indels 66; Gaps 14;

Qy 128 RQIYGYDSRFSGFKDFLLNYPFSTSVKL-STG-----CTGTLVAEKHVLTAACHIDGK 181
      |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||:
Db 28 RVVGGEDARPN-----SNPQWVSQLYDVSSGQRHTCGGTLDVQSWSLVTAAHCISRR 79

Qy 182 TY--VKGTQKURVGFLPKPKFDGGRANDSTAMPEQMFKQWIRVTRTHVPKGWIKGNAN 239
      |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||:
Db 80 TYRVLVGRHSL-----STNEPGSLA-----VKVSKLVVHQDW---NSN 114

Qy 240 DIGNDYDVALLEKKP-HKRKPMKIGVSPPAKQ-LPG-----GRIHFSGVDND--R 286
      |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||:
Db 115 QLSNGNDIALUKLASPVSLTKIQIGCLUPAAQTILLNNVCYYTGWGRLQTNGASPDILQ 174

Qy 287 PGNIVRYRFCVDVKDETYDILLYQQCAQPGAGSAGGVYVRM 324
      |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||:
Db 175 QGQLL-----VVD-----YATC-SKPGWWSGVTKTNM 200

RESULT 6
T29767
hypothetical protein ZC581.6 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
C;Accession: T29767
R;Waterston, B.; Gattung, S.; Le, T.T.
submitted to the EMBL Data Library, May 1997
A;Description: The sequence of C. elegans cosmid ZC581.
A;Reference number: 220882
A;Accession: T29767
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-522 <WAT>
A;Cross-references: UNIPROT:O01771; EMBL:AF003134; PIDN:AAB54144.1; GSPDB:GN00019; CESP:
A;Experimental source: strain Bristol N2; clone ZC581
C;Genetics:
A;Gene: CESP:ZC581.6
A;Map position: 1
A;Introns: 36/2; 138/3; 234/2; 311/1; 331/1; 421/1; 470/2

Query Match          5.6%; Score 117; DB 2; Length 522;
Best Local Similarity 20.4%; Pred. No. 0.082;
Matches 95; Conservative 56; Mismatches 153; Indels 162; Gaps 21;

Qy 29 KPWPAYRLPVULPQSTLINLAKPDFGAELVSSS---CGPOCHKGTPLPTVEEAKQ-- 83
      |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||:
Db 26 KPSNKASASLRKKSNSNPKNKTARSVSKSVPKSAIPASTVQKEVPVVEIKKEEK 85

Qy 84 -----YLSYETLYANGSRRTQVGIIYLSSSGDGAQHRSDGSSGSKRRKRQIYGVDSRF 137
      |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||:
Db 86 PENOKKELAERKL---DRTDQDGKEYEAESALGVVIKDKAPAKMDGDGYDFGPCECF 141

Qy 138 SI FGKO-----FLNYPFSTSV-----KLTGCTGTGLVAEK 168
      |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||: |||:
Db 142 PFFLKILEMLIYWLRIFPSAKVYNGRADSQSEAPWSVFITYLISKDQSATTCGTIVSPR 201

```

QY 169 HVLTAHCI-----HDGK-----TYVKGTKL---RVGFLKPKFKDGGGA 206
Db 202 HILIATCFAGQNRDGSWNLIETDFRSNCKDDYVITNQEFKLKRVFELSNK-----253
QY 207 NDSTSAMPEQMKQWIRVKETHVPKGIKGNANDIGMDY---DYALLEL-----252
Db 254 -KGISRYPEKITLVHACTKKT-----ANRTKKIPQYYTDDFAIVHLVYEELTFSSNVQ 305
QY 253 -----KKPHKR---KFMKIGVSPAKQLPGGRIFHFGYDNDPQNLVY-----292
Db 306 SVCVADETPQNDKLSLEYFGFLNPPSD-----INQNGVDN--TGOLRYEKIEVFRSH 357
QY 293 -----RFGVDKDETVD-----LLYQOCDAQ-----PGASGSGVYVVMKROQKQWE 333
Db 358 PMEIYFFQARDITDKTVACVSVLSKILILNKTOASLNISLKGDSGGGAIDV-----KKG 411
QY 334 RKLIIGIFS-----GHQVDMWNGSPQDFNVAVRITPLKYAQIC 370
Db 412 KTIIGVLSQTSCKRRGGNETMELYSVGPKNQI-----CKYTGIC 453
RESULT 7
B26823
pancreatic elastase II (EC 3.4.21.71) A precursor - human
C:Species: Homo sapiens (man)
C>Date: 16-Aug-1988 #sequence_revision 16-Aug-1988 #text_change 09-Jul-2004
C:Accession: B26823; A27432; A41431; S34491
R:Kawashima, I.; Tani, T.; Shimoda, K.; Takiguchi, Y.
DNA 6, 163-172, 1987
A:Title: Characterization of pancreatic elastase II cDNAs: two elastase II mRNAs are expressed in the rat pancreas
A:Reference number: A90958; MUID:87217962; PMID:3646943
A:Accession: B26823
A:Molecule type: mRNA
A:Residues: 1-269 <RAW>
A:Cross-references: UNIPROT:P08217; GB:M16652; NID:g182057; PIDN:AAA52380.1; PID:g182058
R:Fletcher, T.S.; Shen, W.F.; Largman, C.
Biochemistry 26, 7256-7261, 1987
A:Title: Primary structure of human pancreatic elastase 2 determined by sequence analysis of complementary DNA
A:Reference number: A27432; MUID:88107669; PMID:3427074
A:Accession: A27432
A:Molecule type: mRNA
A:Residues: 1-269 <FILE>
A:Cross-references: GB:M16631; NID:g182022; PIDN:AAA52374.1; PID:g182023
R:Shirasu, Y.; Yoshida, H.; Matsuki, S.; Takemura, K.; Ikeda, N.; Shimada, Y.; Ozawa, T.
J. Biochem. 102, 1555-1563, 1987
A:Title: Molecular cloning and expression in Escherichia coli of a cDNA encoding human pancreatic elastase II
A:Reference number: A41431; MUID:88198076; PMID:2834346
A:Accession: A41431
A:Molecule type: mRNA
A:Residues: 1-201,'V',203-269 <SHI>
A:Cross-references: GB:D00236; NID:g219619; PIDN:BAA00165.1; PID:g219620
A:Note: the authors translated the codon GTG for residue 202 as Cys
R:Moulard, M.; Michon, T.; Kerfelec, B.; Chapus, C.
FEBS Lett. 261, 179-183, 1990
A:Title: Further studies on the human pancreatic binary complexes involving procarboxypeptidase A
A:Reference number: S08253; MUID:90169111; PMID:2307232
A:Accession: S34491
A:Molecule type: protein
A:Residues: 'X',18-50 <MOU>
C:Genetics:
A:Gene: GDB:ELAI
A:Cross-references: GDB:119866; OMIM:130120
A:Map position: 12pter-12qter
C:Superfamily: trypsin; trypsin homology
C:Keywords: hydrolase; pancreas; serine proteinase
F:1-16/Domain: signal sequence #status predicted <SIG>
F:17-28/Domain: propeptide #status predicted <PRO>
F:29-269/Product: pancreatic elastase IIA #status predicted <MAT>
F:29-262/Domain: trypsin homology <TRY>
F:73,121,216/Active site: His, Asp, Ser #status predicted
Query Match 5.6%; Score 115.5; DB 2; Length 269;

Best Local Similarity 28.6%; Pred. No. 0.048;
Matches 50; Conservative 21; Mismatches 51; Indels 53; Gaps 11;
QY 147 NYPFSTSVKLSGTG-----CTGTLVAEKHVLTAACHIDGKTYVKGTKLQKRVGFLKPKFK 200
Db 39 SWPQWVSLQSYSSNGKWHYHTCGGSLTANSWVLTAAHCISSTRY-----RVGL-----85
QY 201 DGGRG-----ANDSTSAMPEQMKQWIRVKETHVPKGIKGNANDIGMDYDYLLELKKP- 255
Db 86 --GRNLVYVAESGSLA-----VSVSKIIVHKDW-----NSNQISKGNIDIALKLANPV 131
QY 256 HKRKWKIGVSPAKQ-LPG-----GRIHFGYDND--RPGNLV---YRFC 295
Db 132 SLTDKIQLACLPPAGTILPNNYPCVYTGWRLQTNAGVDPVLQGGRLLVVDYATC 186
RESULT 8
A25528
pancreatic elastase II (EC 3.4.21.71) precursor - mouse
C:Species: Mus musculus (house mouse)
C>Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 09-Jul-2004
C:Accession: A25528
R:Stevenson, B.J.; Hagenbuechle, O.; Wellauer, P.K.
Nucleic Acids Res. 14, 8307-8330, 1986
A:Title: Sequence organisation and transcriptional regulation of the mouse elastase II ar
A:Reference number: A93646; MUID:87066713; PMID:3641189
A:Accession: A25528
A:Molecule type: mRNA
A:Residues: 1-271 <STE>
A:Cross-references: UNIPROT:P05208; GB:X04573; NID:g50825; PIDN:CAA28242.1; PID:g50826
C:Superfamily: trypsin; trypsin homology
C:Keywords: hydrolase; serine proteinase
F:1-30/Domain: signal sequence #status predicted <SIG>
F:31-271/Product: pancreatic elastase II #status predicted <MAT>
F:31-264/Domain: trypsin homology <TRY>
F:75,123,218/Active site: His, Asp, Ser #status predicted
Query Match 5.6%; Score 115.5; DB 2; Length 271;
Best Local Similarity 25.0%; Pred. No. 0.048;
Matches 67; Conservative 38; Mismatches 80; Indels 83; Gaps 18;
QY 148 YPFSTSVK-LSTG-----CTGTLVAEKHVLTAACHIDGKTY--VKGTKLQKRVGFLKPKFK 199
Db 42 WPMQVSLQVLSGGRWRHNGCGSLVANNWVLTAAHCLSNYQTYRVLLGAHSL-----92
QY 200 KDGGRGANDSTSAMPEQMKQWIRVKETHVPKGIKGNANDIGMDYDYLLELKKP-HKR 258
Db 93 --SNPGAGSAA-----VQVSKLVVHQRW---NSQNVGNGVDIALIKLASPVTL 136
QY 259 KPMKIGVSPAKQ-LPGGRI-HFGY-----DNDPQNLV---YRFC-----295
Db 137 KNIQIACLPPAGTILPRNVYCVYTGWGLLQTNAGSPDTLRQGRLLVVDYATCSSASWMS 196
QY 296 DYKDETY-----DLLYQOCDAQPGASGSGVYVVMKROQKWKERKIIGIFSGHQWVDMNGS 351
Db 197 SYKSSMVCAGGSGVTSSCN-----GDSGPLNCRASNGQ---WQ--VHGIVSFGSSLGCN-Y 247
QY 352 PDENVAVRITPLKYAQICWIKGNVLD 379
Db 248 PRKPSVFRVS-----NYID 262
RESULT 9
S70439
pancreatic elastase I (allele HEU1-16) probable splice form I - human
C:Species: Homo sapiens (man)
C>Date: 03-Aug-1998 #sequence_revision 03-Aug-1998 #text_change 28-Apr-2003
C:Accession: S70439
R:Kawashima, I.; Tani, T.; Mita-Honjo, K.; Shimoda-Takano, K.; Ohmine, T.; Furukawa, H.;
DNA Seq. 2, 303-312, 1992
A:Title: Genomic organization of the human homologue of the rat pancreatic elastase I ger
A:Reference number: A56615; MUID:92338395; PMID:1633328
A:Accession: S70439

A;Description: Isolation, sequencing and characterization of 2 cDNA clones coding for tr

RESULT 13

DBHU
complement factor D (EC 3.4.21.46) precursor [validated] - human (fragment)
N;Alternate names: adipsin; C3 convertase activator
C;Species: Homo sapiens (man)
C;Date: 28-Aug-1985 #sequence_revision 31-Dec-1992 #text_change 09-Jul-2004
C;Accession: A40197; A00936; A60571; S66645
R;White, R.T.; Damm, D.; Hancock, N.; Rosen, B.S.; Lowell, B.B.; Usher, P.; Flier, J.S.;
J. Biol. Chem. 267, 9210-9213, 1992
A;Title: Human adipsin is identical to complement factor D and is expressed at high level
A;Reference number: A40197; MUID:92250520; PMID:1374388
A;Accession: A40197
A;Molecule type: mRNA
A;Residues: 1-246 <WHI>
A;Cross-references: UNIPROT:P00746; GB:M84526
R;Niemann, M.A.; Bhowm, A.S.; Bennett, J.C.; Volanakis, J.E.
Biochemistry 23, 2482-2486, 1984
A;Title: Amino acid sequence of human D of the alternative complement pathway.
A;Reference number: A00936; MUID:85000441; PMID:6383466
A;Accession: A00936
A;Molecule type: protein
A;Residues: 19-44, 'G', 46-51, 'O', 53-75, 'TH', 'P', 80-83, 'XXXITIS', 90-172, 86-91, 185-235, '
A;Note: a few residues were assigned from the previously published sequence of Reid et al
R;Miyata, T.; Oda, O.; Inagi, R.; Sugiyama, S.; Miyama, A.; Maeda, K.; Nakashima, I.; Ya
Mol. Immunol. 27, 637-644, 1990
A;Title: Molecular and functional identification and purification of complement component
A;Reference number: A60571; MUID:90370044; PMID:2395435
A;Accession: A60571
A;Molecule type: protein
A;Residues: 19-20, 'XX', 23-27, 'XX', 30-31, 'XX', 34, 'X', 36-40 <MIY>
R;Baik, N.; Holtkamp, U.; Hoerl, W.H.; Tschesche, H.
FEBS Lett. 371, 300-302, 1995
A;Title: Inhibition of degradation of human polymorphonuclear leukocytes by complement
A;Reference number: S66645; MUID:96013156; PMID:7556615
A;Accession: S66645
A;Status: preliminary
A;Molecule type: protein
A;Residues: 19-44, 'C', 46-48 <BAL>
C;Comment: Factor D cleaves factor B when the latter is complexed with factor C3b, activ
C;Genetics:
A;Gene: GDB:DF
A;Cross-references: GDB:132645; OMIM:134350
A;Map position: Xpter-Xqter
C;Superfamily: trypsin; trypsin homology
C;Keywords: complement alternate pathway; hydrolase; plasma; serine proteinase
F;1-18/Domain: signal sequence #status predicted <SIG>
F;19-246/Product: complement factor D (fragment) #status experimental <MAT>
F;19-241/Domain: trypsin homology <TRY>
F;44-60,141-207,172-188,197-222/Disulfide bonds: #status predicted
F;59,105,201/Active site: His, Asp, Ser #status predicted

Query Match 5.5%; Score 114; DB 1; Length 246;
Best Local Similarity 28.4%; Pred. No. 0.057;
Matches 44; Conservative 20; Mismatches 43; Indels 48; Gaps 9;

QY 149 PFSTSVKLSGTG--CTGTVAEKHLVTAACHIH---DGKTYVKGTKLRVGLFKPKFDGG 203
DB 31 PYMASVQLNGAHLCAGVILVAERVLSAAHCLDAADGKGVQL----- 72
QY 204 RGANDSTSAMPEQMKFWIRVKRTHVPKGIKNANDIGMDYDYLLELKKPKRPMKI 263
DB 73 LGAHSLSQPFPSKRLVDLRA---VP---HPDSQPDTHDHLILLQLSE-----KA 117
QY 264 GVSPPAKQLPGGRTHFGSYDND--RPNGLVYRFCDV 297
DB 118 TLGPAVRPLFWQRV-----DRDVAPFL-----CDV 143

RESULT 14

ELPG
pancreatic elastase (EC 3.4.21.36) I precursor - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 24-Apr-1984 #sequence_revision 30-Sep-1990 #text_change 09-Jul-2004

C;Accession: JS0013; A26777; A10061; A00959
R;Shirasu, Y.; Yoshida, H.; Mikayama, T.; Matsuki, S.; Tanaka, J.I.; Ikenaga, H.
J. Biochem. 99, 1707-1712, 1986
A;Title: Isolation and expression in Escherichia coli of a cDNA clone encoding porcine p
A;Reference number: A92005; MUID:86304235; PMID:3528137
A;Accession: JS0013
A;Molecule type: mRNA
A;Residues: 1-266 <SHI>
A;Cross-references: UNIPROT:P00772; GB:X04036; GB:D00070; GB:N00070; NID:g1941; PIDN:CAA;
R;Tani, T.; Kawashima, I.; Furukawa, H.; Ohmine, T.; Takiguchi, Y.
J. Biochem. 101, 591-599, 1987
A;Title: Characterization of a silent gene for human pancreatic elastase I: structure of
A;Reference number: A26777; MUID:87250343; PMID:3648024
A;Accession: A26777
A;Molecule type: mRNA
A;Residues: 1-125, 'G', 127-183, 'L', 185-266 <TAN>
A;Cross-references: GB:D00160; NID:9217683; PIDN:BA001118.1; PID:g217684
A;Note: the authors translated the codon GGG for residue 58 as Gln, GGC for residue 126 &
R;Shotton, D.M.; Hartley, B.S.
Biochem. J. 131, 643-675, 1973
A;Title: Evidence for the amino acid sequence of porcine pancreatic elastase.
A;Reference number: A90267; MUID:73229121; PMID:4578945
A;Accession: A10061
A;Molecule type: protein
A;Residues: 27-91, 'N', 93-203, 'N', 205-266 <SHO>
R;Shotton, D.M.; Hartley, B.S.
Nature 225, 811-816, 1970
A;Title: Three-dimensional structure of tosyl-elastase.
A;Reference number: A93160; MUID:70114044; PMID:5415110
A;Contents: annotation; X-ray crystallography, 3.5 angstroms; active site
C;Superfamily: trypsin; trypsin homology
C;Keywords: hydrolase; pancreas; serine proteinase; zymogen
F;1-16/Domain: signal sequence #status predicted <SIG>
F;17-26/Domain: activation peptide #status predicted <APT>
F;27-266/Product: elastase I #status experimental <MAT>
F;27-259/Domain: trypsin homology <TRY>
F;56-72,153-220,184-200,210-240/Disulfide bonds: #status experimental
F;71,119,214/Active site: His, Asp, Ser #status experimental

Query Match 5.4%; Score 113; DB 1; Length 266;
Best Local Similarity 27.3%; Pred. No. 0.076;
Matches 36; Conservative 21; Mismatches 41; Indels 34; Gaps 6;

QY 147 NYPFSTSVKLSGTG-----CTGTVAEKHLVTAACHIDGKTY--VKGTQKLRVGLFKPK 198
DB 37 SNFSQISLQYRSGSSWAHTCGGTLRQNWMTAAHCVDRELTFRVVVGHNL----- 88
QY 199 FKDGGERGANDSTSAMPEQMKFWIRVKRTHVPKGIKNANDIGMDYDYLLEL-KKPHK 257
DB 89 -----NQNDGTE-----QYVGQKIVVHPYV---NTDDVAAGYDIALRLAQSRTL 131

QY 258 RKFMKIGVSPPA 269
DB 132 NSYVQLGVLPRA 143

RESULT 15

ELRT1

pancreatic elastase (EC 3.4.21.36) I precursor - rat

C;Species: Rattus norvegicus (Norway rat)

C;Date: 18-Aug-1982 #sequence_revision 18-Aug-1982 #text_change 09-Jul-2004

C;Accession: A00960; A20534

R;MacDonald, R.J.; Swift, G.H.; Quinto, C.; Swain, W.; Pictet, R.L.; Nikovits, W.; Rutte;

Biochemistry 21, 1453-1463, 1982

A;Title: Primary structure of two distinct rat pancreatic preproelastases determined by

A;Reference number: A00960; MUID:82182867; PMID:6918221

A;Accession: A00960

A;Molecule type: mRNA

A;Residues: 1-266 <MAC>

A;Cross-references: UNIPROT:P00773; GB:V01234; NID:g56088; PIDN:CAA24544.1; PID:g56089

R;Largman, C.

Biochemistry 22, 3763-3770, 1983

A;Title: Isolation and characterization of rat pancreatic elastase.

A;Reference number: A20534; MUID:84000385; PMID:6555050

A;Accession: A20534

A;Molecule type: protein

A;Residues: 17-37,'X',39-45 <LAR>

C;Superfamily: trypsin; trypsin homology

C;Keywords: hydrolase; pancreas; serine proteinase; zymogen

F;1-16/Domain: signal sequence #status predicted <SIG>

F;17-26/Domain: activation peptide #status predicted <APT>

F;27-266/Product: elastase I #status predicted <MPT>

F;27-259/Domain: trypsin homology <TRY>

F;71,119,214/Active site: His, Asp, Ser #status predicted

Query Match 5.4%; Score 113; DB 1; Length 266;

Best Local Similarity 26.7%; Pred.No. 0.076;

Matches 39; Conservative 23; Mismatches 48; Indels 36; Gaps 7;

QY 128 RQYGYDSRFSIFGKDFLLNYPFSTSVKLSGT-----CTGTLVAEKHVLTAACHIDGKT 182

Db 26 RVVGGAEARRNSWPSQISLQY-----LSGGSWYHTCGGTLIIRNWMVMTAAHCVSQMT 78

QY 183 YVKGTOKLRVGFLLKPKFKDGRGANDSTSAMPEQMKFQWIRVKETHYKGIKGNANDIG 242

Db 79 F-----RVVVG-----DHNLSQDGT-----QYVSVQKIMVHPTW---NSNVA 115

QY 243 MDYDYALLEL-KKPKKFKFMKIGVSP 267

Db 116 AGYDIALRLAQSVTLNNYVQLVLP 141

Search completed: July 1, 2005, 21:09:20

Job time : 21.3479 secs

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: July 1, 2005, 20:53:37 ; Search time 90.581 Seconds

(without alignments)

2165.204 Million cell updates/sec

Title: US-09-658-677-18

Perfect score: 2080

Sequence: 1 MAGIPGLLFLFLCAVGO.....LKVAICYWIKGNVLDREG 383

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1612378 seqs, 512079187 residues

Total number of hits satisfying chosen parameters: 1612378

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : UniProt 03:*

1: uniprot_sprot:*

2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	2080	100.0	383	1 PS23 HUMAN	O95084 homo sapien
2	1903	91.5	383	2 O6A561	Q9ay61 rattus norv
3	1888.5	90.8	382	1 PS23 MOUSE	Q9d6x6 mus musculu
4	1882.5	90.5	382	2 Q8B2S4	Q8bz84 mus musculu
5	1042	50.1	413	2 Q9BQP6	Q9bqp6 homo sapien
6	1041	50.0	413	2 Q8N320	Q8n320 homo sapien
7	1016.5	48.9	409	2 Q8C0F9	Q8c0f9 mus musculu
8	1014.5	48.8	418	2 Q8GML6	Q8gm16 brachydanio
9	1011.5	48.6	409	2 Q8C0L5	Q8c0l5 mus musculu
10	1005.5	48.3	409	2 Q8C0D6	Q8c0d6 mus musculu
11	194	9.3	316	1 GSEP BACLI	P80057 bacillus li
12	194	9.3	316	2 Q65NR6	Q65nr6 bacillus li
13	188	9.0	450	2 Q71Y85	Q71y85 listeria mo
14	178.5	8.6	364	2 Q98G17	Q98g17 rhizobium l
15	158	7.6	271	2 Q987W6	Q987w6 rhizobium l
16	142.5	6.9	358	2 Q931E7	Q931e7 staphylococ
17	141.5	6.8	321	2 Q7NGB4	Q7ngb4 gloeobacter
18	138.5	6.7	323	2 Q73D54	Q73d54 bacillus ce
19	135.5	6.5	313	1 MPR BACSU	P39790 bacillus su
20	133.5	6.4	799	2 Q6P994	Q6pf94 mus musculu
21	133.5	6.4	811	1 TMS6 MOUSE	Q9db10 mus musculu
22	132.5	6.4	290	2 Q81HL5	Q81hl5 bacillus ce
23	131	6.3	218	2 Q8DR41	Q8dr41 streptococc
24	131	6.3	267	2 Q6DGM4	Q6dgm4 brachydanio
25	129.5	6.2	303	2 Q9EXR9	Q9exr9 bacillus in
26	129	6.2	266	2 Q6AZC0	Q6azc0 brachydanio
27	129	6.2	469	2 Q9GMD9	Q9gmd9 ornithorhyn
28	127.5	6.1	1322	2 Q9NAT0	Q9nat0 anopheles g
29	126.5	6.1	678	2 Q9JJS8	Q9jjs8 rattus norv
30	124.5	6.0	1234	2 Q7P1Q7	Q7p1q7 anopheles g
31	124.5	6.0	1322	2 Q7PNR7	Q7pnr7 anopheles g

32	124.5	6.0	1322	2	Q9NJS5	O9nj55 anopheles g
33	124	6.0	266	1	EL1 BOVIN	O28153 bos taurus
34	124	6.0	286	2	O46644	O46644 macaca fasc
35	123.5	5.9	339	2	Q9QX91	Q9qx91 rattus norv
36	123.5	5.9	366	2	Q9QX85	Q9qx85 rattus norv
37	123.5	5.9	541	2	Q9QX90	Q9qx90 rattus norv
38	123.5	5.9	623	2	Q9JJP3	O9jjp3 rattus norv
39	123.5	5.9	643	2	Q9QX84	Q9qx84 rattus norv
40	122	5.9	259	2	Q9XY61	Q9xy61 ctenecephal
41	122	5.9	376	1	FA10 HOPST	P83370 hoptosephal
42	122	5.9	449	2	Q9VDU8	O9vdus drosophila
43	121.5	5.8	490	1	FA10 RABIT	O19045 cryotolagus
44	121	5.8	745	2	Q90WF9	O90wf9 triakis scy
45	121	5.8	1059	2	Q7Z411	Q7z411 homo sapien

ALIGNMENTS

RESULT 1
PS23 HUMAN STANDARD; PRT; 383 AA.
AC O95084;
DT 28-FEB-2003 (Rel. 41, Created)
DT 28-FEB-2003 (Rel. 41, Last sequence update)
DT 25-OCT-2004 (Rel. 45, Last annotation update)
DE Serine protease 23 precursor (EC 3.4.21.-) (Putative secreted protein
DE ZSIG13) (UNQ270/PRO307).
GN Name:PRSS23; Synonyms=ZSIG13;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Euthera; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RS SEQUENCE FROM N.A.
RC TISSUE=Umbilical vein;
RA Li X., Tedder T.F.;
RT "A novel serine protease from human umbilical vein endothelial
RT cells.";
RL Submitted (JUL-1997) to the EMBL/GenBank/DBJ databases.
RN [2]
RS SEQUENCE FROM N.A.
RA Sheppard P., Blumberg H., Jelinek L., Foster D., O'Hara P.;
RL Submitted (OCT-1999) to the EMBL/GenBank/DBJ databases.
RN [3]
RS SEQUENCE FROM N.A.
RA MEDLINE=21154917; PubMed=11230166; DOI=10.1101/gr.154701;
RC TISSUE=Uterus;
RA Wiemann S., Weil B., Wellenreuther R., Gassenhuber J., Glassl S.,
Ansoerge W., Boecher M., Bloeker H., Bauersachs S., Blum H.,
Lauber J., Duesterhoeft A., Beyer A., Koehler K., Strack N.,
Mewes H.-W., Oettersweil B., Obermaier B., Tampe J., Heubner D.,
Wambutt R., Korn B., Klein M., Pouatka A.;
RA "Towards a catalog of human genes and proteins: sequencing and
RT analysis of 500 novel complete protein coding human cDNAs.";
RL Genome Res. 11:422-435 (2001).
RN [4]
RS SEQUENCE FROM N.A.
RA MEDLINE=22887296; PubMed=12975309; DOI=10.1101/gr.1293003;
RC Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,
Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,
Eaton D., Foster J., Grimaldi C., Gu Q., Hass P.E., Heidens S.,
Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,
Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,
Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,
Vandlen R., Watanabe C., Wieand D., Woods K., Xie M.-H., Yansura D.,
Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,
Godowski P., Gray A.;
RA "The secreted protein discovery initiative (SPDI), a large-scale
RT effort to identify novel human secreted and transmembrane proteins: a
RL bioinformatics assessment.";
RN Genome Res. 13:2265-2270 (2003).
RN [5]

RP SEQUENCE FROM N.A.
 RC TISSUE=Cervix;
 RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
 RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
 RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
 RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
 RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
 RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
 RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullaly S.J.,
 RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
 RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
 RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
 RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
 RA Blakesley R.W., Touchman J.W., Green E.D., Shevchenko Y., Bouffard G.G.,
 RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
 RA Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
 RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
 RT "Generation and initial analysis of more than 15,000 full-length human
 RT and mouse cDNA sequences.";
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
 CC -1- SUBCELLULAR LOCATION: Secreted (Potential).
 CC -1- SIMILARITY: Belongs to the peptidase S1 family.
 CC -----
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 CC -----
 DR EMBL; AF015287; AAD01553.1; -
 DR EMBL; AF193611; AAR07186.1; -
 DR EMBL; ALJ36914; CAB66848.1; -
 DR EMBL; AY359033; AAQ89392.1; -
 DR EMBL; BC001278; AAH01278.1; -
 DR MEROPS; S01.309; -
 DR H-InvDB; HIX0010006; -
 DR InterPro; IPR009003; Pept_Ser_Cys.
 DR InterPro; IPR001254; Peptidase_S1.
 DR InterPro; IPR001314; Peptidase_S1A.
 DR Pfam; PF00089; Trypsin; 1.
 DR PRINTS; PR00722; CHYMOTRYPSIN.
 DR SMART; SM00020; Tryp_SPC; 1.
 DR PROSITE; PS00240; TRYPSIN_DOM; FALSE_NEG.
 DR PROSITE; PS00134; TRYPSIN_HIS; 1.
 DR PROSITE; PS00135; TRYPSIN_SER; FALSE_NEG.
 DR Hydrolase; Serine protease; Signal.
 FT SIGNAL 1 23 Potential.
 FT CHAIN 24 383 Serine protease 23.
 FT ACT_SITE 175 175 Charge relay system (By similarity).
 FT ACT_SITE 240 240 Charge relay system (By similarity).
 FT ACT_SITE 316 316 Charge relay system (By similarity).
 FT DISULFID 160 176 By similarity.
 FT CARBOHYD 93 93 N-linked (GlcNAc...) (Potential).
 FT CARBOHYD 207 207 N-linked (GlcNAc...) (Potential).
 FT SEQUENCE 383 AA; 43001 MW; 46EB6C1ABFD5E8F CRC64;
 Query Match 100.0%; Score 2080; DB 1; Length 383;
 Best Local Similarity 100.0%; Pred. No. 6.2e-168;
 Matches 383; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MAGTPLLFLFLLCAVGQSPYSAPWKTWPAYRLPVVLPOSTLNLAKPFGAEAKLE 60
 DB 1 MAGTPLLFLFLLCAVGQSPYSAPWKTWPAYRLPVVLPOSTLNLAKPFGAEAKLE 60
 QY 61 VSSCGPQCHKGTPLPYEEAKQVLSYETLYANGSRRTQVGVIYILSSSGDGAQRDSGS 120
 DB 61 VSSCGPQCHKGTPLPYEEAKQVLSYETLYANGSRRTQVGVIYILSSSGDGAQRDSGS 120

QY 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLVSTGCTGTLVAEKHVLTAACHIDG 180
 DB 121 SGKSRKRQIYGYDSRFSIFGKDFLLNYPFSTSVKLVSTGCTGTLVAEKHVLTAACHIDG 180
 QY 181 KTVVKGTKLRYGFLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKETHVPKGIKGNAND 240
 DB 181 KTVVKGTKLRYGFLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKETHVPKGIKGNAND 240
 QY 241 IGMVDYDYLLELKKPKHKKFPMKIGVSPAPKQLPGGRIHFSGYDNDPRGNLVYRFCVDKDE 300
 DB 241 IGMVDYDYLLELKKPKHKKFPMKIGVSPAPKQLPGGRIHFSGYDNDPRGNLVYRFCVDKDE 300
 QY 301 TYDLYLQQCDAOPGASGSGVYVMWKRQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
 DB 301 TYDLYLQQCDAOPGASGSGVYVMWKRQOQKWERKIIGIFSGHQWDMNGSPQDFNVAVR 360
 QY 361 ITPLKYAQICYWKGNLYDCREG 383
 DB 361 ITPLKYAQICYWKGNLYDCREG 383
 RESULT 2
 Q6AY61 PRELIMINARY; PRT; 383 AA.
 ID G6AY61;
 AC G6AY61;
 DT 25-OCT-2004 (TrEMBLrel. 28, Created)
 DT 25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
 DT 25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
 DE Hypothetical protein.
 OS Rattus norvegicus (Rat).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
 ON NCBI_TaxID=10116;
 RX [1]
 RP SEQUENCE FROM N.A.
 RC TISSUE=Kidney;
 RC PubMed=12477932; DOI=10.1073/pnas.242603899;
 RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
 RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
 RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
 RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
 RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
 RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
 RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullaly S.J.,
 RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
 RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
 RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
 RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
 RA Blakesley R.W., Touchman J.W., Green E.D., Shevchenko Y., Bouffard G.G.,
 RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
 RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
 Jones S.J., Marra M.A.;
 RT "Generation and initial analysis of more than 15,000 full-length human
 RT and mouse cDNA sequences.";
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
 RN [2]
 RP SEQUENCE FROM N.A.
 RC TISSUE=Kidney;
 RC Director MGC Project;
 RL Submitted (AUG-2004) to the EMBL/GenBank/DBJ databases.
 CC -1- SIMILARITY: Belongs to peptidase family S1.
 DR EMBL; BC079179; AAH79179.1; -
 DR GO; GO:0000786; C:nucleosome; IEA.
 DR GO; GO:0005634; C:nucleus; IEA.
 DR GO; GO:004263; F:chymotrypsin activity; IEA.
 DR GO; GO:0003677; F:DNA binding; IEA.
 DR GO; GO:0007001; P:chromosome organization and biogenesis (gen. .; IEA.
 DR GO; GO:0006334; P:nucleosome assembly; IEA.
 DR GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
 DR InterPro; IPR001951; Histone_H4.
 DR InterPro; IPR001254; Peptidase_S1.

DR InterPro; IPR001314; Peptidase_SIA.
 DR InterPro; IPR009003; Pept_Ser_Cys.
 DR Pfam; PF00089; Trypsin_1.
 DR PRINTS; PR00722; CHYMOTRYPSIN.
 DR ProDom; PD001827; Histone_H4; 1.
 DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.
 KW Hydrolase; Hypothetical protein; Protease; Serine protease.
 SQ SEQUENCE 383 AA; 43159 MW; 4AB12CC7B66CDFC8 CRC64;

Query Match 91.5%; Score 1903; DB 2; Length 383;
 Best Local Similarity 91.4%; Pred. No. 6.4e-153;
 Matches 350; Conservative 11; Mismatches 22; Indels 0; Gaps 0;

QY 1 MAGIPGLLELLFLLLCAGVGSPVSPAPKPTWPAAYRLPVVLPOSTLNLAQDFGAEAKLE 60
 DB 1 MAGIPGLLELLLVLLCVFMQVSPNVPKPTWPAAYRLPVVLPOSTLNLAQDFGAEAKLE 60

QY 61 VSSCGPQCHKGTPLPYEEAKQVLSYETLYANGSRTEQTQVGIYLSNGEGRARSDEA 120
 DB 61 VSSCGPQCHKGTPLPYEEAKQVLSYETLYANGSRTEQTQVGIYLSNGEGRARSDEA 120

QY 121 SGKSRKQIYGVDSRISFGKDFLLNYPSTSVKLTGTCTGLTVABKHVLTAAHCHIDG 180
 DB 121 AGKSRKQIYGVDSRISFGKDFLLNYPSTSVKLTGTCTGLTVABKHVLTAAHCHIDG 180

QY 181 KTVYKGTQKLRVGLKPKKDGGRGANDSTSAMPEQMKFQWIRKTRTHVPKGIKGNAND 240
 DB 181 KTVYKGTQKLRVGLKPKKDGGRGANDSTSAMPEQMKFQWIRKTRTHVPKGIKGNAND 240

QY 241 IGMDDYVALLLEKPKHKKFMKIGVSPPAKQLPGRIHFSGYDNDPRGNLVYRFDVKDE 300
 DB 241 IGMDDYVALLLEKPKHKKFMKIGVSPPAKQLPGRIHFSGYDNDPRGNLVYRFDVKDE 300

QY 301 TYDLLYQQCDAQPGASGSGVYVWVKRQKQKWKRIIGFSGHWDVMDGSPQDFNVAVR 360
 DB 301 TYDLLYQQCDAQPGASGSGVYVWVKRQKQKWKRIIGFSGHWDVMDGSPQDFNVAVR 360

QY 361 ITPKVAQICVWIKGNVLDREG 383
 DB 361 ITPKVAQICVWIKGNVLDREG 383

RESULT 3

PS23_MOUSE STANDARD; PRT; 382 AA.

AC Q9D6X6; Q8V6G1;
 DT 28-FEB-2003 (Rel. 41, Created)
 DT 28-FEB-2003 (Rel. 41, Last sequence update)
 DE 25-OCT-2004 (Rel. 45, Last annotation update)
 DE Serine protease 23 precursor (EC 3.4.21.-).
 GN Name=Prs23;
 OS Mus musculus (Mouse).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Mammalia; Eutheria; Rodentia; Sciurognath; Muridae; Murinae; Mus.
 OX NCBI_TaxID=10090;
 RN [1]
 RP SEQUENCE FROM N.A.
 RC STRAIN=C57BL/6J; TISSUE=Tongue;
 RX MEDLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;
 RA Okazaki Y., Furuno M., Kagukawa T., Adachi J., Bono H., Kondo S.,
 RA Nikaido I., Oeato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,
 RA Yagi K., Tonaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori T.,
 RA Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
 RA Schraml L.M., Kanapin A., Matsuda H., Batalov S., Baisel K.W.,
 RA Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.B., Cousins S.,
 RA Dalla E., Dragani T.A., Fletcher C.F., Forrest A., Frazer J.S.,
 RA Gaasterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,
 RA Grimmond S., Gustincich S., Hirokawa N., Jackson I.J., Jarvis E.D.,
 RA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
 RA Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
 RA Maglott D.R., Maltais K., Marchionni L., McKenzie L., Miki H.,
 RA Nagashima T., Numata K., Okido T., Pavan W.J., Perteau G., Pesole G.,
 RA Petrovsky N., Pillai R., Pontius J.U., Qi D., Ramachandran S.,

Ravasi T., Reed J.C., Reed D.J., Reid J., Ring B.Z., Ringwald M.,
 Sandelin A., Schneider C., Semple C.A., Setou M., Shimada K.,
 Sultana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M.,
 Verardo R., Wagner L., Wahlestedt C., Wang Y., Watanabe Y., Wells C.,
 Wilming L.G., Wynshaw-Boris A., Yanagisawa M., Yang I., Yang L.,
 Yuan Z., Zavolan M., Zhu Y., Zimmer A., Carninci P., Hayatsu N.,
 Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato K.,
 Shiraki T., Waki K., Kawai J., Aizawa K., Arakawa T., Fukuda S.,
 Hara A., Hashizume M., Inotani K., Ishii Y., Itoh M., Kagawa I.,
 Miyazaki A., Sakai K., Sasaki D., Shibata K., Shingawa A.,
 Yasunishi A., Yoshino M., Waterston R., Lander E.S., Rogers J.,
 Birney E., Hayashizaki Y.
 "Analysis of the mouse transcriptome based on functional annotation of
 60,770 full-length cDNAs";
 Nature 420:563-573 (2002).
 [2]

SEQUENCE FROM N.A.
 RP TISSUE=Breast tumor;
 RC MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
 RX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
 RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
 RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
 RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heife F.,
 RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
 RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
 RA Brownstein M.J., Uedlin T.B., Toehiyuki S., Carninci P., Prange C.,
 RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
 RA Bosak S.A., McSwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
 RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
 RA Villalon D.K., Munz D.M., Sodergren E.J., Lu X., Gibbs R.A.,
 RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
 RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
 RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
 RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
 RA Butterfield Y.S.N., Krzywinski M.I., Skalek U., Smailus D.E.,
 RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
 RT "Generation and initial analysis of more than 15,000 full-length human
 and mouse cDNA sequences";
 Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
 RL CC -!- SUBCELLULAR LOCATION: Secreted (Potential).
 CC -!- SIMILARITY: Belongs to the peptidase S1 family.
 CC -----
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 CC or send an email to license@ebi.ac.uk).
 CC -----
 DR EMBL; AK009947; BAB26541.1; -;
 DR EMBL; AK009947; BAB26541.1; -;
 DR EMBL; BC018517; AAH18517.1; -;
 DR HSP; P00746; IDSU.
 DR MEROPS; S01.309; -;
 DR MGD; MGI:1523703; 2310046G1SRIK.
 DR InterPro; IPR009003; Pept_Ser_Cys.
 DR InterPro; IPR001254; Peptidase_S1.
 DR InterPro; IPR001314; Peptidase_S1A.
 DR Pfam; PF00089; Trypsin; 1.
 DR PRINTS; PR00722; CHYMOTRYPSIN.
 DR SMART; SM00020; TRYP_SPC; 1.
 DR PROSITE; PS00240; TRYPSIN_DOM; FALSE_NEG.
 DR PROSITE; PS00134; TRYPSIN_HIS; 1.
 DR PROSITE; PS00135; TRYPSIN_SER; FALSE_NEG.
 KW Hydrolase; Serine protease; Signal.
 FT SIGNAL 1 22 Potential.
 FT CHAIN 23 382 Serine protease 23.
 FT ACT_SITE 174 174 Charge relay system (By similarity).
 FT ACT_SITE 239 239 Charge relay system (By similarity).
 FT ACT_SITE 315 315 Charge relay system (By similarity).
 FT DISULFID 159 175 By similarity.
 FT CARBOHYD 92 92 N-linked (GlcNAc...) (Potential).

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FT CARBOHYD 206 206 N-linked (GlcNAc...) (Potential).
FT CONFLICT 260 260 M -> I (in Ref. 1; BAB26541).
SQ SEQUENCE 382 AA; 43071 MW; 6F09A5C80A5B2306 CRC64;

Query Match 90.8%; Score 1888.5; DB 1; Length 382;
Best Local Similarity 90.8%; Pred. No. 1.1e-151;
Matches 348; Conservative 14; Mismatches 20; Indels 1; Gaps 1;

Qy 1 MAGIPGLLFLFLFLLCAVGVSPYSAPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 60
Db 1 MAGIPG-LFILLVLLCVFMQVSPYTPWKPTWPAYRLPVVLPQSTLNLAKEPFGAEAKLE 59

Qy 61 VSSSCGQCHKGTPLPYEEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRS DGS 120
Db 60 VSSSCGQCHKGTPLPYEEAKQYLSYETLYANGSRTEQVGIYILSSSGDGAQHRS DSEA 119

Qy 121 SGKSRKROIYGYDVSRSIFGKDFLLNYPSTSVKLSCTGCTGLVAEKHVLTAHCIDHG 180
Db 120 TGRSRKROIYGYDGRSIFGKDFLLNYPSTSVKLSCTGCTGLVAEKHVLTAHCIDHG 179

Qy 181 KTYVKGTKLVRGVFLKPKFKDGGRGANDSTSAMPEQMKFQWIRVKTHTVPGWIKGNAND 240
Db 180 KTYVKGTKLVRGVFLKPKFKDGGRGANDSTSSSSAMPDKMKFQWIRVKTHTVPGWIKGNAND 239

Qy 241 IGMVDYVALLLELKKPHKPKFKMGKIVSPAPKQLPGRIHFGSYNDRPGNLVYRFDVKDE 300
Db 240 IGMVDYVALLLELKKPHKPKFKMGKIVSPAPKQLPGRIHFGSYNDRPGNLVYRFDVKDE 299

Qy 301 TYDLYOCCDAQPCASGSGVYVWVKXQOQKWKIIGIFSGHWDVMDGSPQDENVAVR 360
Db 300 TYDLYOCCDAQPCASGSGVYVWVKXQOQKWKIIGIFSGHWDVMDGSPQDENVAVR 359

Qy 361 ITPLYKQAQICVWIKGNLYDCREG 383
Db 360 ITPLYKQAQICVWIKGNLYDCREG 382

RESULT 4
Q8BZS4 PRELIMINARY; PRT; 382 AA.
AC Q8BZS4
DT 01-MAR-2003 (TrEMBLrel. 23, Created)
DT 01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
DE Mus musculus adult male cecum cDNA, RIKEN full-length enriched
DE library, clone:9130215B18 product:SERINE PROTEASE (HYPOTHEICAL 43.0
DE kDa PROTEIN) (PROTEASE, SERINE, 23) homolog.
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Cecum;
RX MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
RA Carninci P., Hayashizaki Y.;
RT "High-efficiency full-length cDNA cloning.";
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Cecum;
RX MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;
RA RIKEN FANTOM Consortium;
RT "Functional annotation of a full-length mouse cDNA collection.";
RN [3]
RP SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Cecum;
RA The FANTOM Consortium;
RA the RIKEN Genome Exploration Research Group Phase I & II Team;
RT "Analysis of the mouse transcriptome based on functional annotation of
RT 60,770 full-length cDNAs.";
RL Nature 420:563-573(2002).
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Qy	241	IGMDYDYVALLLEKPKHKEFKMGIVSPPAKQLPGGRIHPSGYDNDRPGNLVYRFGDVKDE	300
Db	240	IGMDYDYVALLLEKPKHKEFKMGIVSPPAKQLPGGRIHPSGYDNDRPGNLVYRFGDVKDE	299
Qy	301	TYDLLYQOCDAQPGASGSGVYVRMWKROQKWERKIIGIFSGHQWDMNGSPQDFNVAVR	360
Db	300	TYDLLYQOCDAQPGASGSGVYVRMWKROQKWERKIIGIFSGHQWDMNGSPQDFNVAVR	359
Qy	361	ITPLKYAQICYWIKNYLDLCREG 383	
Db	360	ITPLKYAQICYWIKNYLDLCREG 382	
RESULT 5			
Q9BQP6	ID	Q9BQP6	PRELIMINARY; PRT; 413 AA.
AC	Q9BQP6		
DT	01-JUN-2001	(TRENBLrel. 17, Created)	
DT	01-JUN-2001	(TRENBLrel. 17, Last sequence update)	
DT	01-MAR-2004	(TRENBLrel. 26, Last annotation update)	
DE	DJ223E3.1	(Putative secreted protein ZS1313)	
GN	Name=DJ223E3.1;		
OS	Homo sapiens (Human)		
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;		
OC	Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.		
NCBI_TaxID=9606;			
RN	[1]		
RX	SEQUENCE FROM N.A..		
RP	Dunn M.;		
RA	Submitted (MAR-2001) to the EMBL/GenBank/DBJ databases.		
RL	-1- SIMILARITY: Belongs to peptidase family S1.		
CC	EMBL; ALI21939; CAC35071.1; --		
DR	MEROPS; S01.994; --		
DR	Genew; HGNC:21387; PRSS35.		
DR	GO; GO:0004263; F:chymotrypsin activity; IEA.		
DR	GO; GO:0008233; F:peptidase activity; IEA.		
DR	GO; GO:0004295; F:trypsin activity; IEA.		
DR	GO; GO:0006508; P:proteolysis and peptidolysis; IEA.		
DR	InterPro; IPR001254; Peptidase S1.		
DR	InterPro; IPR001314; Peptidase S1A.		
DR	InterPro; IPR003003; Pept_Ser_Cys.		
DR	Pfam; PF00089; Trypsin; 1.		
DR	PRINTS; PR00722; CHYMOTRYPSIN.		
DR	SMART; SM00020; Tryp_SPC; 1.		
DR	PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN_1.		
KW	Hydrolase; Protease; Serine protease.		
QY	SEQUENCE 413 AA; 47098 MW; 818D9C951BD2D6C1 CRC64;		
Query Match 50.1%; Score 1042; DB 2; Length 413;			
Best Local Similarity 47.8%; Pred. No. 7.7e-80;			
Matches 200; Conservative 68; Mismatches 100; Indels 50; Gaps 7			
Qy	7	LELLFLLLCNAV--QVSPYSAPKPTWPAYRLPVLPQSLTINLAKPDPGAELKLVSSS 64	
Db	5	LLMLIFP---TPGWTLDGSEMEWDFMHLRKVPRIVSERTFHLTSPAFADAKQMVNTV 61	
Qy	65	CGPOCHKGTPLPTVEEAKQVLSYETLYANGSRKTETOVGI--VILSSSGDGAQHRDSSGS 122	
Db	62	CGTECQELFPSPSELEDYLSYETVENGRTLTRVKVQDLVEPT-----QNIITKG 115	
Qy	123	KS-RRKROIYGYDSRPSIFGKDFLLNYPFSTSVKLSTGCTGTLVAERHVLTAACHIDHGK 181	
Db	116	VSVRRKQVYGTDSRPSILDKRFLTNPPFSTAVKLSGCSGLILSPQHVLTAACHVHDGK 175	
Qy	182	TYVKGTKLVRGFLKPKPKDGR-----GANDSTSAMPEQMK----- 218	
Db	176	DYVKGSKLRLVGLLMKRNKSGCKRRGSKRSRRREASGGDQREGTREHLRERAKGGRRRK 235	
Qy	219	-----FQWIRVRKTHVPKGIWKNANDIGMDYDVALLEKPKHKEFKMGIV 265	
Db	236	SGRQRTAEGRPPSQWTRVKVNTHTPKGWARGMGDATLDYDVALLEKRAHKHKTWEIGI 295	
Qy	266	SPPAKQLPGGRIHPSGYDNDRPGNLVYRFGDVKDETYDLLYQOCDAQPGASGSGVYVRMW 325	

Submitted (JUN-2004) to the EMBL/GenBank/DBJ databases.
-1- SIMILARITY: Belongs to peptidase family S1.

EMBL; AK031411; BAC27392.1; -
EMBL; BC075675; AAH75675.1; -
MGI; MGI:2444800; Prse35.
GO; GO:0005615; C:extracellular space; TAS.
InterPro; IPR001254; Peptidase_S1.
InterPro; IPR001314; Peptidase_S1A.
InterPro; IPR009003; Pept_Ser_Cys.
Pfam; PF00089; Trypsin; 1.
PRINTS; PR00722; CHYMOTRYPSIN.
SMART; SM0020; Tryp_Spc; 1.
PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN 1.
KW Hydrolyase; Protease; Serine protease.
SEQUENCE 409 AA; 45787 MW; 5E22D4A908E7EF5 CRC64;

Query Match 48.9%; Score 1016.5; DB 2; Length 409;

Best Local Similarity 49.4%; Pred. No. 1.1e-77;
Matches 193; Conservative 68; Mismatches 83; Indels 47; Gaps 5;

QY 31 TWPAYRLPVLPVLPSTLNLAKEPFGAEAKLEVSQCGPOCHKGTPLTYEAKQYLSYETL 90
DB 28 TWLSRIPOVVSNTIHLASPTFOADGVVVKATVCGIECOELPAPSLQLESLSYETI 87
QY 91 YANGSRTEQVGIYL-----SSSGDAQHRDSSGSKRRKQIYGVDSRFSIFGKDF 144
DB 88 FENGTFILRVKQGLVLETRNSVKGAK-----PRRRQVIGTDSRFSILDKRF 138
QY 145 LNLNPFSTSVKLTGCTGTVAEKHVLTAACHDGVKQYVKGQKLRVGLKPKFKDG-- 202
DB 139 ATNFPNTAVKLSTGCGTLVSPNHVLTAAHCHVDGKDYVKGSKLVRVGLKRNKGGK 198
QY 203 -GRGA-----NDSTSAMPEQM-----KFWIRVTRTHVPKG 232
DB 199 KRGSKRSRREASAGSOAHLRESTQRFEGKSRGPRVTRQGRPSQWTRVSKTHIPKG 258
QY 233 WIGNANDIGMDYDYLALLELKKPKKFKWIGVSPPAKQLPGGRIHFGSYDNDRPGNLVY 292
DB 259 WVGNGGLALDYDYLALLELKKRAHQHMLGVSPIITKLPGRRIHFGSFDNRDLSQVY 318
QY 293 RFDVKTEDYLLYQCDADPAGSAGSGVYVVMKQKQKWKRIIGIFSGHGWDMNGSP 352
DB 319 RFCSVSEESNDLLYQYCDAAEGSTGSIYLRLEKPGQKNKRIKIVAVYSQHWVDHVQ 378
QY 353 QDNVAVRITPLKYAQICWIKGNYLDCREG 383
DB 379 KDYNVAVRITPLKYAQICLWIHGNAANCAYG 409

RESULT 8

Q6GML6
ID Q6GML6 PRELIMINARY; PRT; 418 AA.
AC Q6GML6;
DT 05-JUL-2004 (TEMBLrel. 27, Created)
DT 05-JUL-2004 (TEMBLrel. 27, Last sequence update)
DT 05-JUL-2004 (TEMBLrel. 27, Last annotation update)
DE 29c:91804.
GN ORFNames=29c:91804;
OS Brachydanio rerio (Zebrafish) (Danio rerio).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
OC Cyprinidae; Danio.
OX NCBI_TaxID=7955;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Whole;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Schetz T.E.,

Brownstein M.J., Udwin T.B., Toshiyuki S., Carninci P., Prange C.,
Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullaby S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
FAhey J., Helton E., Kettaman M., Madan A., Rodrigues S., Sanchez A.,
Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywiniski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Maizra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]
RP SEQUENCE FROM N.A.
RC TISSUE=Whole;
RA Strausberg R.;
RL Submitted (JUN-2004) to the EMBL/GenBank/DBJ databases.
DR EMBL; BC074028; AAH74028.1; -; 29c:91804.
DR ZFIN; ZDB-GENE-040704-55; zgc:91804.
DR GO; GO:0008233; F:peptidase activity; IEA.
DR GO; GO:0004295; F:trypsin activity; IEA.
DR GO; GO:0006508; P:proteolysis and peptidolysis; IEA.
DR InterPro; IPR001254; Peptidase_S1.
DR InterPro; IPR009003; Pept_Ser_Cys.
DR Pfam; PF00089; Trypsin; 1.
DR PROSITE; PS00134; TRYPSIN_HIS; UNKNOWN 1.
KW Hydrolyase; Protease; Serine protease.
SQ SEQUENCE 418 AA; 47270 MW; AF53345CEB94F720 CRC64;

Query Match 48.8%; Score 1014.5; DB 2; Length 418;

Best Local Similarity 46.6%; Pred. No. 1.7e-77;
Matches 196; Conservative 64; Mismatches 118; Indels 43; Gaps 2;

QY 1 MAGIPGLLELLFLLCAVGOVSPYSAPWKPPTPAVRLPVLPSTLNLAKEPFGAKLE 60
DB 1 MGPVPLTLLLSIALAVLSTTVDPTGDTYTPQKIPLVQEKQTVHLSSEFLAKPQND 60
QY 61 VSSSCQPCQCHKGTPLTYEAKQYLSYETLYANGSRTEQVGIYLLSSSGDGAQRDGS 120
DB 61 LHGICGIECQQRLLPEPSLDLLEQLLSYETMYDNGTRTLTTVTVDLNVSDMT---GAS 116
QY 121 SKSRKRQIYGVDSRFSIFGKDFLNLNPFSTSVKLTGCTGTVAEKHVLTAACHD 180
DB 117 QLHTRHREYGTDTFTITDKQYSMKYPFSTSVKISTGCSGLVSPKPHVLTAAHCH 176
QY 181 KTYVKGTKLRVGLKP-----KFKD 201
DB 177 TDYDGVQKLSVGLKERSRRKRGKRGKRGKRGKRGKRGKRGKRGKRGKRGKRGK 236
QY 202 GGRGANDSTSAMPEQMKFQWIRVTRTHVPKGIKGNANDIGMDYDYLALLELKKPKR 261
DB 237 KGRNRNRSTDEQSPFRWTRVQKQVPGFKGISENVLYDYVAVLELKEAQTKEP 296
QY 262 KIVGSPPAKQLPGRIHFGSYDNDRPGNLVYRCDVKDTEYLLYQCDADPAGSAGSGVY 321
DB 297 DLGVIPSVKLLPAGRIHFGSFDNRDLPGLNLYRFSVSEESNDLLYQYCDAPGSGSGVY 356
QY 322 VVMKQKQKWKRIIGIFSGHGWDMNGSPDENVAVRITPLKYAQICWIKGNYLDCR 381
DB 357 IRLKEPGKKKKWKRIIGIFSGHGWDMNGSQDQYDYNVAVRITPLKYAQICRWVHDS 416
QY 382 E 382
DB 417 D 417

RESULT 9

Q8COL5
ID Q8COL5 PRELIMINARY; PRT; 409 AA.
AC Q8COL5;
DT 01-MAR-2003 (TEMBLrel. 23, Created)


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Db      169  YGSCSATLIGRTVLTAAHCLYSHEDKDWL--SEYLFV-----PGLUNGSTA---- 212
Qy      215  EQMKFQIWRVVKRTHVPKGIWKGNAIDIG--MDYDYALLEKKPKRKFEMKIGVSPPAQOL 272
Db      213  DDAPFGAFTYESAVLQGFIDNYQGYGVSIPWDLGIITLKQDVGTNLGWLGYA-NYDDL 271
Qy      273  PGGRIHSGYNDRPP-GNLVYRPFCDVKDETVDLLYQQ--CDAQPGASGSGVYVRWVKRQQ 329
Db      272  GDFTANLVGYPGDKPMGTMKASCSEVHAENIAPFYQYDCDCTFPGSSGSSVYAYDTKSKQ 331
Qy      330  QKWERKIIGFSGHQWDMNGSPDENVAVRITPLKYAQICVWIKGNY 377
Db      332  R-----IITG--VNVAESPP-DANTAVRLN-----AANVQWINSLY 363

RESULT 15
Q987W6
ID Q987W6 PRELIMINARY; PRT; 271 AA.
AC Q987W6;
DT 01-OCT-2001 (TrEMBLrel. 18, Created)
DT 01-OCT-2001 (TrEMBLrel. 18, Last sequence update)
DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)

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OS Rhizobium loti (Mesorhizobium loti).
OC Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
OC Phyllobacteriaceae; Mesorhizobium.
OC NCBI_TaxID=381;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=MAFF303099;
RC MEDLINE=1082930; PubMed=11214968;
RA Kaneko T., Nakamura Y., Sato S., Asamizu E., Kato T., Sasamoto S.,
RA Nakatani A., Idegawa K., Ishikawa A., Kawashima K., Kimura T.,
RA Kishida Y., Kiyokawa C., Kohara M., Matsuoka M., Matsuoka A.,
RA Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Sugimoto M.,
RA Takeuchi C., Yamada M., Tabata S.;
RT "Complete genome structure of the nitrogen-fixing symbiotic bacterium
RT Mesorhizobium loti.";
RL DNA Res. 7:331-338(2000).
CC -1- SIMILARITY: Belongs to peptidase family S1.
DR EMBL: AP003010; BAB53084.1; -;
DR GO: GO:0004263; F:chymotrypsin activity; IEA.
DR GO: GO:0004295; F:trypsin activity; IEA.
DR GO: GO:0006508; P:proteolysis and peptidolysis; IEA.
DR InterPro: IPR001254; Peptidase_S1.
DR InterPro: IPR001314; Peptidase_S1.
DR InterPro: IPR008256; Peptidase_S1B.
DR InterPro: IPR009003; Pept_Ser_Cys.
DR Pfam: PF00089; Trypsin; 1.
DR PRINTS: PR00722; CHYMOTRYPSIN.
DR PRINTS: PR00839; V8PROTEASE.
DR SMART: SM00020; Tryp_SPC; 1.
DR PROSITE: PS00240; TRYPSIN_DOM; 1.
DR PROSITE: PS00134; TRYPSIN_HIS; UNKNOWN_1.
DR Complete protease; Hydrolase; Protease; Serine protease.
SQ SEQUENCE 271 AA; 30385 MW; 37659307C63D0D9C CRC64;

QY	158	IGCIGILVABEKHVLTAAHLHDGKYYVKGJQKLVGF-LAKPKFADGKRGANDSISAMTEQ	211
Db	56	TGCTAPLISPTLLTAACHCITSPIRQGLGPNLAVIRVTP-----GRASRD---ARP--	105
QY	217	MKFQWIRVKRTHV-----PKGWIKGNANDIGMDYDYLLELKKPKRKKFWKIGVSPPA	269
Db	106	--FGWQAKQWVNPYPYRRPSSGL-----HDVGLIELERPPSPSCHFQOLWSPN	152
QY	270	KQ-----LPGGR-IHFGSYDNRDPGNLYVRFCDVKDE-TYDILLYQOCDAPQAGSGGVYVR	323
Db	153	RODLERLNTRLHHISGYPADKPDGTOWEHSERLDRTITEROLFYSVDTCPGHSGAPV----	209

Qy 324 MWKRQOQXWERKIIGI-----FSGHQWVDMNGSPQD-----ENVAVRITP 363
Db 210 -WIHRQOAGPPVVIVAVHTAGPRPHSGGAWGCRPGVPLAPAGLENRGVRLTP 259

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Job time : 91.581 secs